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RESULT 1
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12-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR11508;
                                                                              Claim 1;
                                                                                                                    Use of thioredoxin alone or with for treatment of malignant cells
                                                                                                                                                              WPI; 1991-117511/16.
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                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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(first entry)
                                                                              27; 41pp; English.
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e.g. B lymphocytic leukaemia,
                                                                                                                                    interferon(s)
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Add46716
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Add090877
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designated MP6/Trx.

(Updated on 09-JAN-2003 to

add missing

Matches

Similarity 4; Conserv

Conservative

0

Mismatches

Indels

0;

Gaps

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2e+06;

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RESULT 2
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ID AAWC
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                                         SXS
                                                             CR AAW09052-W09054 are motifs that are present in thioredoxin or enzymes containing thioredoxin domains, e.g. protein-disulphide oxidoreductases. CR These enzymes are used in a method for producing biologically active, CR correctly branched, disulphide-bonded proteins. The method involves CR reacting the corresponding non-branched, reduced protein with a protein-CR disulphide oxidoreductase enzyme that allows correct disulphide bonding, CR in the presence of glutaredoxin and redox buffer (glutathione disulphide) CR with a redox potential of -270 to -140 mV, to give protein with desired CR bacteria (e.g. Escherichia coli) in the correct conformation. Proteins CR such as (pro)insulin, IGF-I, IGF-II, t-PA, growth hormone, Factor VIII, CR orgalistion factors, interleukins, pancreatic mammalian trypsin inhibitors, interferons, rennin, CR prolactin and human alpha-1 trypsin inhibitors can be made in prokaryotic cells in a biologically active form
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Best Local S
Matches 4
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                                                                                                                                                                                                                                                                                                                 Preparing correctly branched proteins contg. di:sulphide linkages - comprises reacting non-branched reduced protein with protein di:sulphide oxido:reductase enzyme in glutaredoxin and redox buffer.
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) HOLMGREN A.
) ASLUND F.
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otic recombinant production; biologically active;
disulphide oxidoreductase; glutaredoxin; redox buffer.
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100.0%; Score 31; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            used in branched protein synthesis method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 31;
Pred. No.
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AAY49348 standard;

peptide; 4 AA

14-MAR-2000 AAY49348;

(first entry)

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                                                                                                                                            The present polypeptide is a conserved motif found in the Trx-homology domain of the thioredoxin (Trx) family of proteins. The motif is important for catalytic activity. The N-terminal region of PKC-cinteracting cousin of thioredoxin (PICOT) shows homology to the Trx-homology domain but it lacks the conserved motif and contains instead an Chial-pro-Gln-Cys motif. PICOT modulates the function of PKCtheta and Cinhibits activation of c-Jun N-terminal kinase (NK) and the Crascription factors Ap-1 and NF-kappaB. These polypeptides are involved in the TCR/CD28 signaling cascade which leads to IL-2 production. PICOT ameliorating physiological conditions, e.g. an inflammation or stress response, associated with these polypeptides. The cDNA encoding PICOT was screened for binding to polypeptides encoded by a Jurkat T lymphoma cDNA colones were sequenced and the putative 37.5 kDa polypeptide encoded by the open reading frame was named PICOT
                                                        Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                      A PKC-interacting cousin of Trx (PICOT) polypeptides and the nucleic acids that encode it, useful e.g. for modulating Jun N-terminal kinase, AP-1 and/or nuclear factor-kappa-B expression and activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB13337 standard; protein; 4 AA
                                                                                                                     Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-442379/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; thioredoxin; Trx; PICOT; PKC-interacting cousin of thioredoxin; protein kinase; Tktei; PKCtheta; c-Jun N-terminal kinase; JNK; AP-1; NFkappaB; interleukin-2; IL-2; TCR/CD28 signaling cascade; PICOT-homology domain; PIH domain; protein activity modulation; stress
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB13337;
                                                                                                                                                                                                                                                                                                                                                                                                                          Example 4; Page 43; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human Trx-homology domain conserved motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (LJOL-) LA JOLLA INST ALLERGY & IMMUNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inflammation.
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                                                                         Similarity
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 2452
                          CGPC 5
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                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99WO-US030285
                                                                     100.0%;
                                                        Score 31; DB 3;
Pred. No. 2e+06;
Mismatches
                                                                                        DB 3;
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                                                                                      Length 4;
                                                          Indels
                                                          0;
                                                          Gaps
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RESULT 4

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XX AAY4
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ABP56403
ID ABP5
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                consisting of a candidate interactor peptide, a conformation-constraining protein and a DNA binding moiety and/or gene activating moiety. The cells are useful for detecting protein interactions. The cells may also be used in a method for identifying and purifying genes encoding a wide range of useful proteins based on their physical interaction with a second
                                                                            CN1189539-A
                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Saccharomyces; mammalian; fusion protein; interactor peptide; conformation-constraining protein; DNA binding moiety; thioredc gene activating moiety; protein interaction; gene purification.
                         05-AUG-1998
                                                                                                                                                                                      Thioredoxin;
                                                                                                                                                                                                                                            Thioredoxin related peptide #2
                                                                                                                                                                                                                                                                                                                                                        ABP56403;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a population of Saccharomyces and/or mammalian cells comprising recombinant DNA molecules encoding fusion proteins, eac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Col 15; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Population of Saccharomyces and/or mammalian cells comprising recombinant DNA encoding fusion proteins, useful for detecting protein interactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-JUL-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-JUL-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                           ABP56403 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GEMY ) GENETICS INST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                      TrxA; fusion protein; allogenic protein.
                                                                                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94US-00278082
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                                                                                                                                                                                                                                                                                                                                                                                                        peptide;
                                                                                                                                                                                                                                                                                                  entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 coli thioredoxin.
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Pred. No. 2e+
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                structure and the DNA sequence of an encoded allogenic protein. The soluble expression of the fusion protein is efficient. The expressed fusion protein can be easily purified as several amino acid sequences inserted at active sites of the thioredoxin gene sequence, and hydrolysed by a specific protease as the junction between the two encoded DNA sequences can be cut by the specific protease, obtaining a recombined allogenic protein the same as the natural protein in one-class structure. The present sequence represents a peptide which is given in the exemplification of the present invention
                                                                                                                                                                                                                           chronic disease; autoimmune disease; asthma; septic shock; lung fibrosis glomerulonephritis; atherosclerosis; autoimmune encephalomyelitis; soft tissue injury; reduction catalyst; chronic inflammatory disease; systemic inflammatory response; NF-kappaB-mediated inflammatory disease; inducible surface expression; CD44; intercellular adhesion molecule; ICAM-1; NF-kappaB activation; redox active peptide.
                                                                                                                                                                                                                                                                                                         nephrotropic; antiarteriosclerotic; antirheumatic; antiarthritic; Helicobacter pylori protein; inhibitor; activator; NF-kappaB; rheumatoid; inflammator; bowel disease; autoimmune arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-JAN-1997;
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 WPI; 2002-627466/67
                           Windle HJ,
                                                                                   30-JAN-2001;
                                                                                                              30-JAN-2002; 2002WO-IE000011
                                                                                                                                                                      WO200260933-A2
                                                                                                                                                                                                                                                                                                                                                     Antiinflammatory; immunosuppressive; antiasthmatic; antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                 Redox active peptide sequence CGPC
                                                                                                                                                                                                                                                                                                                                                                                                                30-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                           AA022593;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAO22593 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               is composed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-733391/80
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                                                                                                                                           08-AUG-2002
                                                                                                                                                                                                  Unidentified
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                                                       (QUEE-) QUEEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expression carrier and its fusion
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention describes a DNA molecule of a fusion protein which of the DNA sequence similar to encoded thioredoxin in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 2 (Disclosure); 14pp;
                           Dermot
                                                                                   2001IE-00000069
                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                         ELIZABETH COLLEGE DUBLIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97CN-00100362
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                           ζ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     peptide; 4
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                           Dermot 0,
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Pred. No. 2e+
0; Mismatches
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                             Abdel-Latif MM,
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2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein of novel thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         activated, autoimmune diseases, asthma, septic shock, lung fibrosis, glomerulonephritis, atherosclerosis or autoimmune encephalomyelitis, and soft tissue injury. The Helicobacter pylori protein is also used for treating chronic diseases where NF-kappaB is transcriptionally activated. The Helicobacter pylori protein catalyses the reduction of proteins. The H. pylori protein is also useful for treating chronic inflammatory diseases, particularly systemic inflammatory response syndrome or other NF-kappaB-mediated inflammatory disease states, down regulates the resting and inducible surface expression of CD44 and intercellular adhesion molecule (ICAM)-1, prevents NF-kappaB activation both prior to stimulation with inducers of NF-kappaB and secondary to induction of NF-kappaB, and prevents or inhibits constitutive NF-kappaB activity. This sequence represents the redox active peptide sequence CGPC relating to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a Helicobacter pylori protein, its derivative fragments, mutants and variants capable of inhibiting the activation NF-kappaB and having a sequence of 106 amino acids defined in the specification. The protein is useful for preventing and/or treating inflammation, inflammatory bowel disease, rheumatoid/autoimmune arthritis, chronic disease where NF-kappaB is transcriptionally
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel Helicobacter pylori protein useful for treating inflammation, autoimmune diseases, septic shock and glomerulonephritis, has redox active peptide sequence inhibiting activation of NF-kappaB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                attenuated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      M. tuberculosis thioredoxin active site peptide
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                                                                                                                                                                                                                                                                                                     07-FEB-2001; 2001US-0267328P.
18-SEP-2001; 2001US-0322989P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-AUG-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mycobacterium tuberculosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tuberculostatic; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-NOV-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABG32509 standard; peptide;
                                                                                         WPI; 2002-643387/69
                                                                                                                                                     Kernodle
                                                                                                                                                                                                                                                                                                                                                                                                 07-FEB-2002; 2002WO-US003451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200262298-A2
                                                                                                                                                                                                               (USGO )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N
                                                                                                                                                                                                                                              UNIV VANDERBILT
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                                                                                                                                                                                                               SD
                                                                                                                                                  DS,
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                                                                                                                                                                                                               DEPT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                intracellular bacterium.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TRX; tuberculosis; antibacterial; active site;
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                                                                                                                                                  Bochan
                                                                                                                                                                                                               VETERANS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             superoxide dismutase; SOD; bacterial infection
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Pred. No. 2e+
0; Mismatches
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he activation of
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Modifying a bacterium to enhance immunogenicity, as vaccines preventing bacterial infections, e.g. tuberculosis comprises

for reducing the

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RESULT 8
AAU97916
X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X D X 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to modifying a bacterium to enhance immunogenicity of the bacterium comprising reducing the activity of an anti-apoptotic enzyme produced by the bacterium e.g. superoxide dismutase (SOD), where the bacterium has enhanced immunogenicity in a subject. Also included are the modified bacterium, an attenuated intracellular bacterium (further modified to reduce the activity of an anti-apoptotic enzyme of the bacterium), modifying a bacterium so it retains or increases immunogenicity but loses or reduces pathogenicity in a subject, comprising reducing but not eliminating an activity of an enzyme produced by the bacterium. Where reducing the activity of the enzyme attenuates the bacterium. The methods are useful for preventing bacterial is infections, e.g. tuberculosis. The attenuated intracellular bacterium is useful as a vaccine for preventing bacterial infections. The present sequence is the active site of M. tuberculosis thioredoxin (TRX) which was eliminated by mutagenesis to create an attenuated bacterium of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antioxidant; oxidative stress; central nervous system; cancer; ageing; neurodegenerative disorder; Parkinson's disease; Alzheimer's disease; Creutzfeldt Jakob's disease; cerebral ischaemia; multiple sclerosis; motor neuron disease; scrapie; spongyform encephalopathy; acute respiratory distress syndrome; amyotrophic lateral sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               activity of an anti-apoptotic enzyme, e.g. superoxide dismutase produced by the bacterium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4 AA;
                                WPI; 2002-419127/45
                                                                                           Atlas D;
                                                                                                                                                                                                                    26-OCT-2000; 2000GB-00026254
                                                                                                                                                                                                                                                                           26-OCT-2000; 2000GB-00026254
                                                                                                                                                                                                                                                                                                                                                                                                          GB2368339-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      atherosclerosis; acquired immunodeficiency syndrome; AIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tetrapeptide antioxidant peptide composition CB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bacterium
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                                                                                                                                                         RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "N terminal acetylation"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94; 164pp; English.
                                                                                                                                                         & DEV CO
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Pred. No. 2e+06;
Mismatches 0;
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disease, motor neuron diseases, scrapie, spongyform encephalopathy and loss or impaired memory, peripheral tissue disorders, particularly acute respiratory distress syndrome, amyotrophic lateral sclerosis, atherosclerotic cardiovascular disease, multiple lateral organ
                                                                                                                                                                                                                                                                                                  may be used for preventing or reducing oxidative stress and for treating central nervous system neurodegenerative disorders, particularly Parkinson's disease, Alzheimer's disease, Creutzfeldt Jakob's disease, cerebral ischaemia, multiple sclerosis, basal ganglia degenerative
                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to an antioxidant compound comprising a peptide including at least 3 amino acid residues of which at least 2 are cysteine residues each having a readily oxidisable sulfhydryl group for effecting antioxidation, and a first hydrophobic or non charged group attached to an amino terminal of the peptide and a second hydrophobic or non charged group attached to a carboxy terminal. the antioxidant peptide compound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antioxidant compound used for treating e.g. Alzheimer's disease and multiple sclerosis comprises peptide including cysteine residues have sulfhydryl groups and hydrophobic or non charged groups attached to
                                                                        dysfunction, complication resulting from inflammatory processes, acquiimmunodeficiency syndrome, AIDS, cancer and ageing. The antioxidant compound of the invention has the advantage that it may cross a blood barrier. The present sequence represents an antioxidant peptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                62pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     having
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RESULT 9
ABB05573
                                                                                                                   Matches
                                                                                                                            Query Match
Best Local Similarity
ABB05573;
                    ABB05573 standard; peptide; 4 AA.
                                                                                             2 CGPC 5
                                                                                                                  4; Conservative
                                                                                                                            100.0%; Score 31; 100.0%; Pred. No.
                                                                                                                   ٥,
                                                                                                                   Mismatches
                                                                                                                            DB 5;
2e+06;
                                                                                                                   0
                                                                                                                                     Length 4;
                                                                                                                   Indels
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                                                                                                                   Gaps
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8

Sequence 4 AA;

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X D X P X P X P X P X P X P X P F F F F E X S X X X X D X X

Modified-site Modified-site

WO200192466-A2

WPI; 2002-114335/15 Rabenstein DA, 26-MAY-2000; 2000US-00579883 16-MAY-2001; 2001WO-US015923

VIND

CALIFORNIA Shi T;

Dithiol; disulfide bond; chemical reagent; sulfhydryl group; SH trans-(Pt(ethylenediamine)2Cl2)2+.

group;

Synthetic.

Location/Qualifiers /note= "acetylated"

Dithiol

peptide

22-APR-2002 (first entry)

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RESULT 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amino acids derivatised for chemical peptide synthesis, where at least cone of the vials comprises an amino acid that, when fully deprotected, bears a SH group, and a vial comprising (1); (3) chemically synthesising a peptide comprising a disulfide linkage, involving chemically coupling a number of amino acids to form a peptide comprising at least two SH groups, and contacting the peptide with (1); and (4) a kit for forming an intramolecular disulfide bond in a peptide, comprises a container disulfide bond in a peptide, comprises a container disulfide bond in a peptide, comprises a container evaluating the efficacy of particular reagent species, and as a sample solution when optimising a purification protocol to clean up or to isolate the desired reaction product (e.g., when optimising a high cressure liquid chromatography (HPLC) protocol or evaluating a particular chromatography column). The present sequence represents a dithiol peptide which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Forming an intramolecular disulfide bond in a peptide, comprises contacting a peptide comprising at least two sulfhydryl groups with an oxidizing agent that forms disulfide linkages in peptides.
New isolated DNA encoding a disulfide bond formation (Dsb) protein bacteria of the genus Ehrlichia, useful for preventing human monocytotropic ehrlichiosis, or canine monocytic ehrlichiosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disulphide bond formation protein; Dsb protein; gene therapy; vaccine; HME; human monocytotropic ehrlichiosis; canine monocytic ehrlichiosis; CME; antibacterial; immunostimulant; thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                comprising a peptide which comprises at least two sulfur groups, and (I) (2) a peptide synthesiser (III) for the synthesis of a peptide having an intramolecular disulfide linkage, comprises a number of vials containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a method of forming an intramolecular disulfide bond in a peptide, comprising contacting a pentide comprisi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E. coli thioredoxin active site peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-AUG-2003
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                                                                                                   WPI; 2003-441356/41.
                                                                                                                                                                                                                                          01-NOV-2001; 2001US-0335611P
                                                                                                                                                                                                                                                                                         01-NOV-2002; 2002WO-US035226
                                                                                                                                                                                                                                                                                                                                                                                        WO2003038061-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                    Escherichia coli.
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                                                                                                                                                                                               (RERE-) RES
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                                                                                                                                                  Walker DH;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and polymucleotides encoding such proteins from bacteria of the genus Ebrlichia. Sequences of the invention are useful for preventing human monocytotropic ehrlichiosis (HME) or canine monocytic chriichiosis (CME) The kits are useful for determining whether an animal or individual has been infected with a species of the bacteria of the Ehrlichia genus. Date sequences are also used in gene therapy and as vaccines. The present sequence is Escherichia coli thioredoxin protein active site peptide (conserved cysteine motif) used in the exemplification of the invention
             their coding sequences, where the thioredoxin molecule is resistant to the oxidizing effects of cytokines or reactive oxygen species or resistant to S-nitrosylation of a SH-group by nitrous oxide. The mutant Trx proteins are useful for treating or diagnosing atherosclerosis, diabetes, apoptotic disease, mitochondrial dysfunction disease or cardiac dysfunction. Mitochondrial dysfunction diseases include Kearns-Sayre
                                                                                                                                                                                              New mutant thioredoxin molecule which is resistant to oxidizing effects of cytokines or reactive oxygen species, useful for treating or diagnosing atherosclerosis, diabetes, Alzheimer's disease, myocardial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antiparkinsonian; Anticonvulsant; Cardiant; Cardiovascular; Antirheumatic; Antiarthritic; Cytostatic; Antipsoriatic; Gethioredoxin; Trx; atherosclerosis; diabetes; apoptotic diss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mitochondrial disease; cardiac dysfunction; angiogenesis dependent disease; neurodegene
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                                                                                                                                                                                             diagnosing
                                                                                                                                                                                                                                                                                              Min W,
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                                                                                                                                                                                                                                                                                                                                                                                             22-JUL-2003; 2003WO-US022847
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 10; Page
                                                                                                                                             Disclosure; SEQ ID NO 15; 145pp;
                                                                                                                                                                                                                                                               WPI; 2004-157111/15.
                                                                                                                                                                                                                                                                                                                              (UYRP)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rheumatoid arthritis; cancer; psoriasis; catalytic site.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Trx catalytic site consensus sequence,
                                                                                                                                                                               infarction
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2004013283-A2
                                                                                               present invention relates to mutant thioredoxin (Trx) ir coding sequences, where the thioredoxin molecule is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CGPC 4
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                                                                                                                                                                                                                                                                                                                              UNIV ROCHESTER
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or encephalomyopathy
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                                                                                                                                                                               or cancer.
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Pred. No. 2e+
0; Mismatches
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                                                                                                                                               English
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No.
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2e+06;
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RESULT 12
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Matches
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             The specification describes a method for increasing the liquefaction of mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum in the patient with a composition comprising a protein or peptide containing a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a total volume of a sample of mucus or sputum from the patient shows a statistically significant increase after administration of the composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian the composition further comprises nicotinamide-adenine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease, Parkinson's disease, Huntington's disease, amytrophic lateral sclerosis, multiple sclerosis or peripheral neuropathy. Cardiac dysfunction includes myocardial infarction, cardiomyopathy, arterial hypertension or heart failure. The mutant Trx proteins are also useful for treating or diagnosing angiogenesis dependent diseases, e.g. rheumatoid arthritis, cancer or psoriasis. The present sequence is the consensus sequence for the catalytic site of the Trx protein (residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Increasing the liquefaction of excessively viscous or cohesive mucus or sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mucus; sputum;
lung disease;
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11-APR-2003; 2003US-0462082P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Amino acid sequence of the thioredoxin active site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-JUN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NAJE-) NAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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form) (NADPH)
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RESULT 13
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                      The present sequence is that of the antioxidant peptide, CB4. This peptide is used in a claimed pharmaceutical composition of the invention also comprising an L-DOPA amide derivative. The pharmaceutical composition is useful for treatment of a condition associated with impaired dopaminergic signalling, such as Parkinson's disease, Tourette' syndrome, attention deficit hyperactive disorder, generation of pituitar tumours and schizophrenia. The inclusion of an antioxidant in the pharmaceutical composition provides for improved stability particularly in vivo, and also provides neuroprotection against oxidative stress, which is typically associated with L-DOPA treatment, and with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thioredoxin active site of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for treating lung diseases (i.e. cystic fibrosis) in which abnormal or excessive viscosity or cohesiveness of mucus or sputum is a symptom or cause of the disease. The present sequence represents an active site of thioredoxin which is used in the method of the invention.
                                                                                                                                                                                                                          New L-dihydroxy phenyl alanine derivatives useful for the treatment of e.g. Parkinson's disease, Tourette's syndrome, attention deficit hyperactive disorder and schizophrenia.
                                                                                                                                                                                               Disclosure; Page 18; 44pp; English.
                                                                                                                                                                                                                                                                                          WPI; 2004-625378/60
                                                                                                                                                                                                                                                                                                                       Atlas
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense therapy; thioredoxin inhibitor; cancer; neoplasm; carcinoma; melanoma; basal cell carcinoma; ovary tumor; breast tumor; comentation; cancer; renal cell carcinoma; bladder tumor; comentatic cancer; panoreas tumor; lung tumor; colorectal tumor; cervical dysplasia; colon tumor; colorectal tumor; cervical dysplasia; colon tumor; colorectal tumor; cardiovascular disease; musculoskeletal disease; dermatological; cardiovascular disease; metabolic disease; neumatoid arthritis; cardiovascular disease; collar disease; rheumatoid arthritis; cardiovascular disease; collar disease; rheumatoid arthritis; colorectal tumor; colorectal tumor; colorectal colorectal; colorectal colorectal colorectal; colorectal c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                       comprise a subsequence of 8 or more nucleotides or nucleotide analogs, wherein the subsequence is located within a sequence selected from ADV11669-ADV11724. The oligonucleotides preferably contain at least nucleotide analog, especially a locked nucleic acid (LNA) or a modifie
                                                                                                                                                                                                                                                                                           Novel compound useful for modulating expression of gene involved in cancer disease, or for modulating red blood cell proliferation, cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-MAR-2005
24-FEB-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADV11816 standard; peptide; 4
                                                                                                                        putative human oncogene
                                                                                                                                            The invention relates to antisense oligonucleotides consisting of 8-50 nucleotides and/or nucleotide analogs which inhibit expression of the
                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                    proliferation,
                                                                                                                                                                                                                                                                                         cancer disease,
                                                                                                                                                                                                                                                                                                                                                                        WPI; 2005-056301/06.
                                                                                                                                                                                                                                                                                                                                                                                                                      Hansen B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-FEB-2003; 2003US-0446374P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2004241717-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cellular organisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Thioredoxin conserved active site motif, SEQ ID NO:149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SANT-) SANTARIS PHARMA AS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-FEB-2004; 2004US-00776933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modulation;
                                                                                                                                                                                                                                                                                                                                                                                                                    Thrue CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A
                                                                                                                                                                                                                      SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                                                                                                                                                                                                                    ion metabolism or glucose and energy metabolism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note = Formation of this disulfide bond is reversible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 31;
100.0%; Pred. No.
                                                                                                                                                                                                                    149; 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cell cycle; thioredoxin; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                 Westergaard
                                                                                                                   thioredoxin (TRX). The antisense oligonucleotides
5-methylcytosine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
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                                                                                                                                                                                                                                                                                                                                                                                                                      Petersen
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                         or a modified
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                                                                                                                                                                                                                                                              gliomas. The TRX antisense oligonucleotides are also useful in the treatment of conditions such as atherosclerosis, psoviasis, diabetic retinopathy, rheumatoid arthritis, asthma, warts, and allergic dermatitis. They may additionally be used for inhibiting cellular proliferation and for modulating red blood cell proliferation, ion metabolism, glucose and energy metabolism, pH regulation, matrix metabolism, apoptosis, cytokinesis or the cell cycle. The TRX antisense oligonucleotides of the invention have increased specificity and affinity for TRX mRNA, and are resistant to degradation. The present sequence represents the reversibly disulfide-bonded conserved active site motif of thioredoxin which is referred to in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chemotherapeutic agent, an antiinflammatory compound or an antiviral compound. The TRX antisense oligonucleotides, and conjugates and compositions containing them, are useful in the treatment of cancers as carcinomas (e.g., malignant melanoma, basal cell carcinoma, ovaria carcinoma, breast carcinoma, non-small cell lung cancer, renal cell carcinoma, bladder carcinoma, recurrent superficial bladder cancer, stomach carcinoma, prostatic carcinoma, pancreatic carcinoma, lung carcinoma, cervical carcinoma, cervical dysplasia, laryngeal papillomatosis, colon carcinoma, colorectal carcinoma and carcinoid tumors); sarcomas (e.g., osteosarcoma, Bwing's sarcoma, chondrosarcom malignant fibrous histiocytoma, fibrosarcoma, and Kaposi's sarcoma);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polynucleotide moieties covalently attached to the oligonucleotide; and a pharmaceutical composition comprising a TRX antisense oligonucleotide or conjugate of the invention, optionally further comprising a champerborance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pseudoisocytosine, 5-bromouracil, 5-propynyluracil, 6-aminopurine, aminopurine, inosine, diaminopurine and 2-chloro-6-aminopurine. The invention also relates to a conjugate comprising a TRX antisense oligonucleotide of the invention and one or more non-nucleotide or
                                                                                                                                                                              Sequence 4 AA;
1 CGPC 4
                                        2 CGPC 5
                                                                                    4; Conservative
                                                                                                            Similarity
                                                                                                                                                                                                                             record
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                                                                                                            100.0%;
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                                                                                                            .0%; Score 31;
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DS Hom
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADW97624 standard; peptide; 4 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-MAR-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP2005060408-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chronic bronchitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human thioredoxin protein-related active center
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                                                                                                                                                                                                                                                                                13-FEB-2004; 2004JP-00071691
                                                                                                                                                                                                                                                                                                                                                                                               30-NOV-2004; 2004JP-00346572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           interleukin 2 antagonist; interleukin 4 antagonist;
interleukin 6 antagonist; TNF alpha antagonist; emp
                                                                                                                                                                        (JPCJ-) JPC KK
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WPI; 2005-200153/21.

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RESULT 16
RDY81685
ID ADY81685
AC ADY81
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention relates to a novel prophylactic or therapeutic agent of chronic obstructive pulmonary disease which contains a polypeptide of thioredoxin super family as active ingredient, preferably human thioredoxin. The invention may be used for the production of compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Agent e.g. tablet or capsules, obstructive pulmonary disease, family as active ingredient.
                                                                                                                         New prokaryotic cell that is status of the cytoplasm to a
                                                                                                                                                                                                                                                                                                                                                                                         05-OCT-1999;
08-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             with a respiratory-Gen. or antiinflammatory activity acting as cytokine-
antagonists, histamine-antagonists, interleukin-2-antagonists,
                                                  Example 6;
                                                                                                  disulfide bond
                                                                                                                                                                                                WPI; 2005-239572/25.
                                                                                                                                                                                                                                           Beckwith J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-OCT-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US6872563-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antiparasitic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genetic engineering;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thioredoxin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADY81685 standard;
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                                                  SEQ ID NO
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                                                                                                                                                                                                                                           Aslund F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000US-00679705
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                                                                                           cytoplasm to
d containing
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99US-0163939P.
99US-0166044P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ering; protein production; antimicrobial; antifungal;
antiviral; immunostimulant; thioredoxin.
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                                                                                                                                                                                                                                                                                                                       COLLEGE
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                                                                                                                      genetically more oxidati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 31;
Pred. No.
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invention relates

to a

prokaryotic cell that is genetically modified

This sequence represents a conserved active site found in thioredoxins. The thioredoxins are a class of low molecular weight redox proteins. Thioredoxin exerts specific redox control over a number of transcription

DNA binding and

therefore regulates

to modulate

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RESULT 17
AAY78911
ID AAY78
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            favors disulfide bond formation, relative to a prokaryotic cell that is not genetically modified. The cell further comprises a mutated AhpC gene comprising an insertion of three nucleotides in the TCT triplet rich region located at codons 36-39 of an AhpC gene. The insertion increases the cell's ability to proliferate relative to a cell that is not further genetically modified. Also described is a method for producing a protein having at least one disulfide bond. The prokaryotic cell and method are useful for producing disulfide bond containing proteins. The proteins, polypeptides, and compositions are used for diagnostic, therapeutic, or prophylactic purposes. They are useful for inhibiting the growth of bacteria, viruses, fungi, or other parasites and for stimulating an immune response. They may be used as food supplements as well as
                                                                                                                                                                                                                                                                                                                                                                                                                                               wound healing; Crohn's disease; ulcerative colitis; angiogenesis; adult respiratory distress syndrome; ARDS; infection; inflammatory conditions, psoriasis; inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             apoptosis; reperfusion injury; hepatitis; amyotrophic lateral scler
Alzheimer's disease; autoimmune disease; Sjogren's syndrome; lupus;
rheumatoid arthritis; HIV; Hermansky-Pudlack syndrome; retinopathy;
                                                                                                                    New composition for preventing inhibition of apoptosis and for treating e.g. cancer or reperfusion injuries comprises inhibitor of cellular redox
                                                                                                                                                                                                                                                                            31-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thioredoxin reductase; redox inhibitor; cellular proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4 AA;
                                                                            Disclosure;
                                                                                                                                                                     WPI; 2000-195158/17.
                                                                                                                                                                                                Kirkpatrick DL,
                                                                                                                                                                                                                                                                                                           02-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                        WO200006088-A2
                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY78911;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY78911 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        components of wash
                                                                                                                                                                                                                                                                                                                                          10-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                  hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             to shift the redox status of the cytoplasm to a more oxidative state that
                                                                                                                                                                                                                               (KIRK/) KIRKPATRICK D
(POWI/) POWIS G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 CGPC 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                          Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              conserved active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                              98US-00127219
                                                                                                                                                                                                                                                                                                           99WO-US017496
                                                                            12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sh powders and creams. The present redox active disulfide motif.
                                                                                                                                                                                                Powis
                                                                                                                                                                                                                                                                                                                                                                                                                                      disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide;
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                                                                          55pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              site amino acid sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                  fungoides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence represents
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sclerosis;
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RESULT 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local
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12-JUL-1991;
19-AUG-1991;
01-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               T cell-derived colony stimulating factor; TC-CSF; immunostimulant; acquired immune deficiency syndrome; AIDS; cancer; gene therapy; haematopoietic growth factor; human; thioredoxin; ADF; active site
The invention relates to T-cell derived colony stimulating factors (TC-CSF) and nucleic acid molecules encoding them. TC-CSF which is a novel haematopoietic growth factor is useful for treating immune-compromised
                                                                                                                                                                                           New T cell-derived colony stimulating factor for treating compromised patients, including acquired immunodeficiency patients and certain types of cancer patients.
                                                                                                                                                                                                                                                                                                                                      WPI; 2001-432041/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transcription.
                                                                                                                                                                                                                                                                                                                                                                                                   Choudhury
                                                                                                                                        Example 33;
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                                                                                                                                 Col 36; 38pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                89US-00356006.
91US-00729135.
91US-00747784.
91US-00788115.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94US-00180371
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -00906866
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B
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                                                                                                     Query Match
Best Local S
Matches 4
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                                   The invention relates to the use of thioredoxin as a cell growth stimulator, for modulating thioredoxin cell growth stimulator-associated inhibition of apoptosis, comprising controlling the expression and/or activity of the thioredoxin. The invention is useful for treating tumours associated with aberrant thioredoxin expression and activity. This sequence represents a wild type thioredoxin active site peptide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            patients, including acquired immune deficiency syndrome (AIDS) patients and certain types of cancer patients. TC-CSF DNA is used in gene therap. The present sequence is human thioredoxin active site which is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cytostatic; thioredoxin; cell growth stimulator; modulating; apoptosis; tumour; wild type.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wild type thioredoxin active site peptide sequence
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                                                                                                                                                                                                              Sequence 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 1; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treatment of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-470971/50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modulating thioredoxin-associated inhibition of apoptosis for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (POWI/) POWIS
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                                                                                                        Conservative
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                                                                                                                                                                                                                                                                     invention
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                                                                                                        100.0%; Score 31; DI
100.0%; Pred. No. 2e-
tive 0; Mismatches
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Pred. No.
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                                                                                                                                    2e+06;
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                                                                                                           Gaps
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ABG71301 ID ABG XX

ABG71301 standard; peptide;

5 84 RESULT 20

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The invention relates to isolated human thioredoxin polypeptide TRXP-2, a naturally occurring polypeptide comprising an amino acid sequence 90% cc identical to TRXP-2, a biologically active or immunogenic fragment of CC TRXP-2 and its encoding polynucleotide (TRXP-2 NA). Also included are a recombinant polynucleotide comprising a promoter sequence operably linked to TRXP-2 NA, a cell transformed with the recombinant sequence, a TRXP-2 CC transgenic organism, an anti-TRXP-2 antibody and a composition comprising CC TRXP-2, an agonist or antagonist compound identified using TRXP-2, or the cabove antibody. TRXP-2 is useful for screening a compound for condition associated with decreased or overexpression of functional TRXP cin a patient. TRXP-2 is useful for screening for a compound that CC modulates the activity of the polypeptide or that binds to the polypeptide. TRXP-2 is also useful as an immunogen for preparing cuseful for screening a compound that condition a compound antibodies by hybridoma technology. TRXP-2 NA is polyclonal or monoclonal antibodies by hybridoma technology. TRXP-2 NA is careful for detecting the presence of TRXP-2 NA. The careful for screening a compound for contibody is useful for detecting the presence of TRXP-2 NA. The careful for detecting the presence of TRXP-2 NA and modulators of composing a condition or disease associated with expression of TRXP in a subject or in a biological sample. TRXP-2 NA and TRXP-2 NA and modulators of conditerative (e.g cancer, actinic keratosis, arteriosclerosis, condition, osteoperation, hepatitis, mixed connective tissue confidence, adult respiratory distress syndrome, Addison's confidence, adult respiratory distress syndrome, Addison's confidence and confidence and prevention or confidence and confidence and antipologism and modulators of confidence and confidenc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inflammatory disorder; acquired immunodeficiency syndrome; infection; adult respiratory distress syndrome; Addison's disease; allergy; asthmatherosclerosis; gout; myocardial inflammation; scleroderma; pericardial inflammation; osteoporosis; rheumatoid arthritis; systemic lupus erythematosus; ulcerative colitis; Reiter's syndrome; haemodialysis; Crohn's disease; atopic dermatitis; viral disorder; autoimmune thyroiditis; diabetes mellitus; Graves' disease; glomerulonephritis; active site disulphide/dithiol conserved sequence
                                           scleroderma, systemic lupus erythematosus, ulcerative colitis, Reiter's syndrome, haemodialysis, Crohn's disease, atopic dermatitis, autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel human thioredoxin preventing and treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (INCY-) INCYTE PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-SEP-2001; 2001US-00954846
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Thioredoxin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein active
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98US-00107248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Guegler KJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein and polynucleotide for diagnosing, cell proliferative, inflammatory and viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   atherosclerosis; bursitis;
and glomerulonephritis)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Baughn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AIDS; trauma;
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RESULT 21
AAP71087
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AC AAP71
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Best Local
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31-OCT-2002
06-JUN-1991
                            It is a redox active peptide which can be used in cpds. to treat stress and ischaemia, for inhibiting lipid peroxidm, due to drugs or toxic effects, as anti-inflammatory agents, to prevent DNA damage, and to maintain the biological activity of antibiotics or proteins. Thioredoxin can be used as general antioxidant in the food and cosmetic industry. It may be used in prode processes for antibiotics, proteins or polypeptides. Therapeutic dose is 1 microgram/kg - 100mg/kg. (Updated on 21-0CT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct
                                                                                                                                                                             New and known thioredoxin derivs. metal catalysed oxidative damage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      infection (e.g. viral, bacterial, fungal, parasitic, protozoal an helminthic), viral disorders (e.g. acquired immunodeficiency synd (AIDS) and trauma. The present sequence represents a thioredoxin site disulphide/dithiol conserved sequence as found in TRXP-2
                                                                                                                                          Claim
                                                                                                                                                                                                                                        Pigiet
                                                                                                                                                                                                                                                                                       14-MAR-1986;
20-OCT-1986;
                                                                                                                                                                                                                                                                                                               10-FEB-1986;
                                                                                                                                                                                                                                                                                                                                      10-FEB-1987;
                                                                                                                                                                                                                                                                                                                                                                16-SEP-1987
                                                                                                                                                                                                                                                                                                                                                                                                                         Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Redox active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence of thioredoxin analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAP71087;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAP71087 standard; peptide; 6
                                                                                                                                                                                                                  WPI; 1987-258442/37.
                                                                                                                                                                                                                                                                                                                                                                                     EP237189-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       food industry;
                                                                                                                                                                                                                                                                (REPK ) REPLIGEN CORP.
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                                                                                                                                         Page 7; 7pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CGPC 5
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milarity 100.
Conservative
                                                                                                                                                                                                                                        Millis CD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     peptide; antioxidant; stress; ischaemia; lipid peroxidation;
y; cosmetics; antibiotic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
                                                                                                                                                                                                                                                                                     86US-00828112.
86US-00839857.
86US-00921287.
                                                                                                                                                                                                                                                                                                                                      87EP-00301150
                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "optional,
terminal blocking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                              /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                          English
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches

    and analogues - useful for preventing
in biological reactions and disease

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Sequence 6

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RESULT 22
ABP54936
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                                                                                                                       The present sequence is that of a hexapeptide comprising the active site CC sequence of thioredoxin. The hexapeptide was used in an example from the cinvention that examined the binding of 4 (N-(S-C)) to peptide and protein CC dithiols. The results indicated that GSAO selectively binds proteins CC containing closely spaced thiols. The invention provides a method for the treatment and/or prophylaxis of arthritis using a compound of formula A-CC (L-Y)p, where A is a cell-membrane impermeable pendant group, L is a CC linker and/or spacer, Y is an arsenoxide or arsenoxide equivalent, and p is an integer from 1 to 10. The compound, e.g. GSAO, is useful for the treatment of arthritic conditions such as calcific periarthritis, CC septic or Jaccoud's arthritis, gout, hand or knee osteoarthritis, conditions from the symovial tissue and leucocyte ingress that triggers inflammation, with no signs or symptoms of toxicity or ingress that triggers inflammation, with no signs or symptoms of toxicity
                                                Query Match
Best Local S
Matches 4
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Best Local S
Matches 4
                                                                                                    Sequence 6
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                                                                                                                                                                                                                                                                                                                                                                                         Use of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-MAR-2001; 2001AU-00003798
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           osteopathic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-MAR-2002; 2002WO-AU000310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200274305-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Active site peptide of thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABP54936;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABP54936 standard; peptide; 6 AA.
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                                                Similarity 4; Conserv
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                        CGPC 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Donoghue N;
                                                                                                     A
A
                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           organoarsenical; arsenoxide; antiarthritic; antirheumatic; arthritis; therapy.
                                                                                                                                                                                                                                                                                                                                                                             vertebrates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
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                                                            100.0%; Score 31;
100.0%; Pred. No.
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Pred. No.
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                                                              2e+06;
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                                                                         DB 5; Length 6;
                                                                                                                                                                                                                                                                                                                                                                                        for the treatment
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                                                Gaps
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RESULT

23

ADH05024
ID ADH
XX
AC ADH
XX
DT 11-I

ADH05024;

ADH05024 standard; peptide; 6 AA

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11-MAR-2004

(first entry)

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RESULT 24
                                                                                                            Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                   The present invention relates to a method for producing an antiangiogenesis plasmin fragment. The method comprises contacting a plasminogen polypeptide with a plasminogen activator and a plasmin reductase, where a reduced plasmin protein is produced and the antiangiogenesis plasmin fragment having anti-angiogenesis activity, is released from the reduced plasmin protein. The plasminogen activator is preferably a urokinase-type plasminogen activator. The angiogenesis plasmin fragment is A61. The plasmin reductase is selected from annexin II heterotetramer, annexin II p36 subunit, p11, thioredoxin, and protein disulphide isomerase. The annexin II heterotetramer is associated with a cell membrane. The method of the invention is useful for modulating (e.g. promoting or inhibiting) angiogenesis. The present sequence represents a mammalian cleavage peptide found in thioredoxin and protein disulphide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  plasmin reductase; reduced plasmin protein; plasminogen activate urokinase-type plasminogen activator; A61; annexin II heterotetramer; annexin II p36 subunit; annexin II p11 subunit; thioredoxin; protein disulphide isomerase; modulating angiogenesis; mammalian; cleavage peptide.
                                                                                                                                                                             Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Producing an anti-angiogenesis plasmin fragment, useful for modulating, e.g. promoting or inhibiting angiogenesis, comprises contacting a plasminogen polypeptide with a plasminogen activator and a plasmin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-596985/56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Waisman D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2003083234-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mammalia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADA08451 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-NOV-2001; 2001US-0333866P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-NOV-2002; 2002US-00304287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (WAIS/) WAISMAN D. (KWON/) KWON M.
                                                                                                              Local Similarity
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                                                                              CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 6;
                                                                                                            100.0%; Score 31; Diarity 100.0%; Pred. No. 2e Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kwon M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      found in thioredoxin and protein disulphide isomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide; 6 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   29pp; English.
                                                                                                                           DB 6;
                                                                                                                                          Length 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      plasminogen activator;
                                                                                                               Indels
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RESULT 25
ADM32929
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a molecule with two pendant phenylarsine moieties. The invention is useful for imparting detectable properties to target material, for detecting target material, for monitoring binding process, for isolating and immobilizing at least one target material, for screening a library of peptides and identifying a cysteine containing peptides and for labeling protein in living cells. The present sequence represents a target peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New bis-phenylarsine derivatives useful for labeling protein in living cells and for screening a library of peptides and identifying a cysteine
                                                                                         Modified-site
                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                   17-JUN-2004
                                                                                                                                                                                                                                                          ADM32929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-JUN-2002; 2002US-0388699P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-JUN-2003; 2003WO-US018792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pendant phenylarsine moieties; labeling; target peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Target peptide
           25-MAR-2004
                                                                                                                                                                                    mucus; sputum; thioredoxin; active site; cohesive mucus; cohesive sputum;
                                                                                                                                                                                                             Amino acid
                                                                                                                                                                                                                                                                                ADM32929 standard; peptide; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ebright RH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003107010-A1
                                  WO2004024868-A2
                                                                   Modified-site
                                                                                                                  Active-site
                                                                                                                                                                         Bunt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RUTF ) UNIV RUTGERS STATE NEW JERSEY
                                                                                                                                                                         disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2004-156348/15.
                                                                                                                                                                                                                                                                                                                                           N
                                                                                                                                                                                                                                                                                                                                                                  2 CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                         4;
                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 5; 82pp; English
                                                                                                                                                                                                           sequence
                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ebright YW,
                                                                                                                                                                                                                                                                                                                                           ហ
                                                                                                                                                                        cystic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             #5 of the invention
                                                                                                                            Location/Qualifiers
                                                        note= "this"
                                                                              note= "this residue is"
                                                                                                     'note= "active site of thioredoxin"
                                                                                                                                                                                                           of the thioredoxin active site
                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 31; DB 8; 100.0%; Pred. No. 2e+06;
                                                                                                                                                                         fibrosis.
                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                        residue is
                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                        in a reduced state
                                                                              in a reduced state"
                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                               Length 6;
                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a tott volume of a sample of mucus or sputum from the patient shows a statistically significant increase after administration of the composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin. The composition further comprises nicotinamide-adenine thioredoxin. The composition further comprises nicotinamide-adenine dinucleotide phosphate (reduced form) (NADPH) for reducing the thioredoxin active site of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or abnormally viscosis or cohesiveness of the liquefaction of excessively or treating lung diseases (i.e. cystic fibrosis) in which abnormal or treating lung diseases (i.e. cystic fibrosis) in which abnormal or the protein of the disease of the disease of the cystic fibrosis) in which abnormal or the protein of the protein an active site of the protein of the protein and the protein of the protein of excessively or cohesiveness of mucus or sputum is a symptom or the protein of the protein an active site of the protein of the protein and the protein of the prot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the patient with a composition comprising a protein or peptide containing a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-SEP-2002;
11-APR-2003;
                       Misc-difference
                                                                                                                                           Synthetic.
                                                                                                                                                                                                                      mucus; sputum; thioredoxin; active site; cohesive mucus; cohesive sputum;
                                                                                                                                                                                                                                                                       Amino acid
                                                                                                                                                                                                                                                                                                                      17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                       ADM32928;
                                                                                                                                                                                                                                                                                                                                                                                                                      ADM32928 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum. The method comprises contacting the mucus or sputum of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 13; Page 51; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Increasing the liquefaction of excessively viscous or cohesive mucus sputum comprises contacting the mucus or sputum with a composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-270016/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-SEP-2003;
                                                                        Active-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     thioredoxin which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprising a protein or peptide containing a thioredoxin active-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (NAJE-) NAT
                                                                                                                                                                                            lung disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CGPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the disease. The present sequence represents an 
xin which is used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CGPC 5
                                                                                                                                                                                                                                                                       sequence of the thioredoxin active site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JEWISH MEDICAL & RES CENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2003WO-US028526
                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002US-0409960P
2003US-0462082P
                                                                                                                                                                                            cystic
                                                                                             Location/Qualifiers
/note= "any amino acid"
                                              note= "active site of thioredoxin"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     describes a method for increasing the liquefaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             represents an active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8; Length 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for
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RESULT 27
ADP87509
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  SX I
                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                         statistically significant increase after administration of the composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian thioredoxin. The composition further comprises nicotinamide-adenine dinucleotide phosphate (reduced form) (NADPH) for reducing the thioredoxin active site of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for the protein active file of the protein and thioredoxin active file of the protein of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for the protein active file of the protein and thioredoxin reductase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mucus or sputum. The method comprises contacting the mucus or sputum of the patient with a composition comprising a protein or peptide containing a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a total volume of a sample of mucus or sputum from the patient shows a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The specification describes a method for increasing the liquefaction of mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum. The method comprises contacting the mucus or sputum of
                                                                                                                                                                                                                                                                                                                                                                    treating lung diseases (i.e. cystic fibrosis) in which abnormal or excessive viscosity or cohesiveness of mucus or sputum is a symptom or cause of the disease. The present sequence represents an active site of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Increasing the liquefaction of excessively viscous or cohesive mucus or sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Misc-difference
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ADP87509;
                                        ADP87509 standard;
                                                                                                                                                                                                                                                                                                          Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 12; Page 51; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           reduced state.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-SEP-2002;
11-APR-2003;
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                                                                                                                                                                                                                                                                                                                                                   thioredoxin which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NAJE-) NAT JEWISH MEDICAL & RES CENT
                                                                                                                                            N
                                                                                                                                                                                   2 CGPC 5
                                                                                                                                                                                                                         4.
                                                                                                                                                                                                                                                Similarity
                                                                                                                                            CGPC
                                                                                                                                                                                                                           Conservative
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2003US-0462082P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "this residue is in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "this residue is in a reduced state"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  note= "any amino acid'
                                                                                                                                                                                                                                                                                                                                                   is used
                                        peptide; 6 AA.
                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                The present sequence represents an active site of a used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          : "any
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino
                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                              Score 31; DB 8;
Pred. No. 2e+06;
                                                                                                                                                                                                                           Mismatches
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D
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                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                             Length 6;
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                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                      cyclohexa-2,5-dienone derivatives (I) and their salts, esters, amides, solvates, hydrates or protected forms. (I) have cytostatic activities, and can be used as thioredoxin reductase inhibitors. (I) can be used for treating a proliferative condition (especially colon or renal cancer) or a condition mediated by thioredoxin/thioredoxin reductase in the human or radiutal body. (I) are also useful for inhibiting thioredoxin/thioredoxin proliferation and/or to promote apoptosis in a cell in vitro or in vivo. The present sequence represents a thioredoxin family conserved catalytic site peptide, which is given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4-(1(Sulfonyl)-1H-indol-2-yl)-4(hydroxy)-cyclohexa-2; 5-dienone;
cytostatic; thioredoxin reductase inhibitor; proliferative condition;
colon cancer; renal cancer; thioredoxin; thioredoxin reductase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New 4-(1(sulfonyl)-1H-indol-2-yl)-4(hydroxy)-cyclohexa-2,5-dienone derivatives are thioredoxin reductase inhibitors useful for treating e.g. colon or renal cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cell cycle progression; cell proliferation; apoptosis; catalytic site peptide.
                                       protein disulfide :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Stevens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-DEC-2002; 2002WO-GB005842
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-SEP-2004
                                                    p11; p11 activity modulator; plasminogen activation; cytostatic;
antisense therapy; cancer; tumour; tumour growth inhibition; thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                  Sequence 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 82; 141pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-517586/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-DEC-2002; 2002WO-GB005842
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO2004056361-A1
                                                                                              Thioredoxin
                                                                                                                          23-SEP-2004
                                                                                                                                                      ADP67423;
                                                                                                                                                                                 ADP67423 standard; peptide; 6 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CANC-) CANCER RES TECHNOLOGY LTD
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                                                                                                                                                                                                                                                                                N
                                                                                                                                                                                                                                                                                                            4.
                                                                                                                                                                                                                                                    CGPC
                                                                                                                                                                                                                                                                                CGPC 5
                                                                                                                                                                                                                                                                                                                                                                   A,
                                                                                              and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention described 4-(1(Sulfonyl)-1H-indol-2-yl)-4(hydroxy)-5-dienone derivatives (I) and their salts, esters, amides,
                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                          (first entry)
                                                                                                                                                                                                                                                      v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Westwell
                                                                                             protein disulfide isomerase common
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          conserved catalytic site peptide
                                                                                                                                                                                                                                                                                                                       100.0%; Score 31; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ₽,
                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Poole TD,
                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wells
                                                                                                                                                                                                                                                                                                                         2e+06;
                                                                                                                                                                                                                                                                                                                                        DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ຸດ
                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                     Length
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                                                                                                                                                                                                                                                                                                             Indels
                                                                                             sequence
                                                                                                                                                                                                                                                                                                                                         0
                                                                                                SEQ
                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                 NO:4.
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Synthetic

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RESULT 29
ADQ97008
ID 7008
ID 70097
AC ADQ97
AC ADQ97
XX 07-00
CT 07-00
XX Thior
XX p11 F
XX p11 F
XX pntl
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                                                                                                                                                                                                                                                                                                                                                          S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a composition (I) which modulates the CC activity of a p11 protein and effects a change in the level of CC plasminogen activation by a cell. Also described: (1) making (M1) a CC clonal cell line, which involves isolating a cell, then characterising CC the activity of a protein produced by the cell or clonal progeny of the CC clonal cell line (II) useful in the identification of composition that CC modulate p11 activity, where the clonal cell line is obtained by (M1). (C) has cytostatic activity, and can be used in antisense therapy. (I) is useful for modulating the activity of p11 which involves administering CC ibinosarcoma cell. The cell can be a human cancer cell, chosen from a HT1080 CC ibinosarcoma cell, a LNCap prostate cancer cell and a CCL-22 colorectal adenocarcinoma cell, a LNCap prostate cancer reducing the development of cancer cell in the patient. (I) is useful for reducing the growth of the modulate p11 activity of a cancer cell invasion in a patient, which involves administering (I) to a cancer cell in the patient. (II) is useful for inhibiting the growth of concert cell in the patient of concert cell in the patient. (II) is useful for concert cell in the patient. (II) is useful for concert cell of a clonal cell line that modulates p11 activity which involves administering the composition that modulates p11 activity which involves cell of a clonal cell line that had not received the composition, and cell interpretation that produces a change in p11 activity. The change in that produces a change in p11 activity. The composition that produces a change in p11 activity. The composition sequence represents a thioredoxin and protein confidentification sequence, which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 4
                                                                                                                                                                      ADQ97008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Composition useful for inhibiting growth of tumor in patient, modulates activity of pl1 protein and effects change in level of plasminogen activation by a cell.
                                         pll protein; plasminogen activation; tumour growth; metastasis;
                                                                                   Thioredoxin
                                                                                                                            07-OCT-2004
                                                                                                                                                                                                             ADQ97008 standard; peptide; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-DEC-2002; 2002US-0433140P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-DEC-2003; 2003WO-US040029
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; SEQ ID NO 4; 140pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-487995/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MEDI-) MEDIOMICS
(WAIS/) WAISMAN D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present
                                                                                                                                                                                                                                                                                                                                                                                                   . Similarity 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIOMICS LLC
                                                                                                                                                                                                                                                                                                                                                          CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AA;
                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 31; DB 8; Length 6; ilarity 100.0%; Pred. No. 2e+06; Conservative 0; Mismatches 0; Indels
                                                                                   and protein disulphide isomerase common
                                                                                                                            (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention.
                                                                                                                            entry)
                                                                                 sequence.
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RESULT 30
AAPS0203
ID AAPS0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a composition which modulates the activity of a pl1 protein and effects a change in the level of plasminogen activation by a cell. The composition is useful for reducing pl1 protein activity in cancer cells. They are also useful for inhibiting tumour growth or metastasis or treating cancer. The present sequence represents the amino acid sequence of the thioredoxin and protein disulphide isomerase common sequence involved with cleavage, formation and reshuffling of disulphide
                                                                                                                                                                                                                                               Sequence of oligopeptide 3 specific to the beta-subunit of luteinising hormone (\operatorname{LH}).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New composition which modulates the activity of a pll protein and effects a change in the level of plasminogen activation by a cell useful for treating cancer or reducing pll protein activity in cancer cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-NOV-2001; 2001US-0333866P
26-NOV-2002; 2002US-00304287
13-DEC-2002; 2002US-0433140P
                                   Maes RF;
                                                                                                                                   22-MAY-1985.
                                                                                                                                                                                                           Contraception; assay; pregnancy test; luteinising hormone; human chorionic gonadotropin.
                                                                                                                                                                                                                                                                                                              AAP50203;
                                                                                                                                                                                                                                                                                                                                     AAP50203 standard; peptide; 8 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; SEQ ID NO 4; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-579971/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Waisman DM;
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                                                                                   26-AUG-1983;
                                                                                                            23-AUG-1984;
                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                     17-OCT-1991
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                                                          (ANDA-)
                                                                                                                                                                                                                                                                                                                                                                                                    2 CGPC 5
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                                                           ANDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                           BIOLOGICALS
                                                                                                                                                                                                                                                                                  (first entry)
                                                                                   83EP-00401714.
                                                                                                            84EP-00401710
                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 31; DB 8;
Pred. No. 2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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WPI; 1985-124152/21

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RESULT 31
ADC81369
ID ADC81369
XX ADC81369
XX COW;
DE BOVI
XX COW;
XW COW;
XW COW;
XW COW;
XW PTI
XW PTI
XW PTI
XW PTI
XX BO
XX PFI
PTI
XX PFI
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Con A-induced interleukin-2 production; mouse spleen cells; granulocyte-monocyte colony forming cell; GPCC; mouse bone marrow; immunomodulatory agent; congenital immune defect; primary T cell deficiency; thymic dysplasia; Down's syndrome; viral infection; herpes; adenovirus infection; HV infection; subacute sclerosing pan-encephalitis; immune suppression; leukopaemia; cancer; chemotherapy; radiotherapy; autoimmune inflammatory disorder; rheumatoid arthritis; systemic lupus erythematosus; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           preparation of anti-HCG and anti-LH vaccines. The vaccines are characterised in that these oligopeptides are complexed upon themselves and between themselves by disulphur bridges, upon their own beta-subunit of HCG and LH or upon oligopeptides containing cysteine whose AA chain is common to LH and HCG, namely the peptides in AAP50208-P50212. Through sulphur bridges the specific oligopeptides are administered by injection in vaccines to cause prodn. of antibodies so that a contraceptive effect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligopeptides AAP50198-P50205, specific to the beta-subunit of LH and HCG, located near cysteine groups, are used in a claimed process for t preparation of anti-HCG and anti-LH vaccines. The vaccines are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prepn. of anti-HCG and anti-LH vaccines - by use of oligo-peptide(s)
specific to beta-sub-unit of LH and HCG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 17; 22pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bovin THF-gamma2 analogue C3-C6-cyclicTHF gamma2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC81369 standard; peptide; 8 AA.
                                       Rycus
                                                                                                       (YEDA ) YEDA RES & DEV CO LTD
                                                                                                                                                                                                                                                  28-JUN-1994;
                                                                                                                                                                                                                                                                                                                      12-JAN-1995
                                                                                                                                                                                                                                                                                                                                                                                      WO9501182-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mutant; mutein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bone marrow transplantation; atopic condition; asthma; atopic dermatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cow; thymic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           achieved; and the body fluids
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                                          Burstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                humoral factor gamma2; THF-gamma2; concanavalin
                                                                                                                                                                                                                                                  94WO-US007304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
3. .6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "Wild-type Lys substituted by Cys"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "Wild-type Asp substituted by Cys"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          note= "This bond cycylises the molecule"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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                                          Trainin N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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14-OCT-1994; 06-OCT-1995; 25-APR-1996. WO9611943-A1

94SE-00003526

95WO-SE001151

(ASTR) ASTRA AB

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein natural or non-natural AA residue, (iv) cyclisation through a free carboxyl group and a free amino group or through disulphide bonds of cysteine residues or (v) linkage of two or more sequences or modified sequences corresponding to (i)-(iv), either directly or through a peptide or non-peptide chain. The peptides are used as immunomodulatory agents. They can be used for treating e.g. congenitel immune defects involving primary T cell deficiencies such as thymic dysplasia and Down's syndrome, primary and secondary viral infections (e.g. herpes virus, adenovirus and HIV), as well as subacute infections used as subacute sclerosing panencephalitis, immune suppression and leukopaenia following cancer treatment by chemotherapy and/or radiotherapy, autoimmune inflammatory disorders, e.g. rheumatoid arthritis, systemic lupus erythematosus and psoriasis, in bone marrow transplantation to prevent viral infections and in atopic conditions such as asthma and atopic dermatitis. The present sequence is a bovine THF-gamma2 analogue peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A peptide is claimed which is a thymic humoral factor (THF)-gamma2 analogue of at least 4 amino acid (AA) residues or a functional derivative or salt, capable of enhancing concanavalin A (Con A)-induced interleukin-2 (IL-2) production in mouse spleen cells and/or the number of granulocyte-monocyte colony forming cells (GM-CFC) of mouse bone marrow. The peptide comprises one or more sequences corresponding to the sequence of the wild-type THF-gamma2 but differing by (i) deletion of or more AA residues at the N-and/or C-terminus, (iii) substitution of one or more AA residues by a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New thymic humoral factor gamma 2 analogues - used as immunomodulatory agents for treating e.g. immune defects, viral infections or auto:immune
                                                                                                                                                                                                                                                                                                       Immunomodulation; immunomodulator; immune system; anergy; deficiency;
aberrant response; chronic bronchitis; atopic disease; AIDS;
acquired immune deficiency syndrome; herpes virus infection; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Page 17; 30pp; English
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                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                    Thymic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW05002 standard;
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                                                                                                                                                                                                           Disulfide-bond
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                                                                                                                                                                                                                                                                                                                                                                                  factor gamma-2 cyclic analogue.
                                                                                                                                                                                                           Location/Qualifiers
3. .6
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Pred. No. 2e+
0; Mismatches
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2e+06;
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RESULT 33
AAW41664
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A novel immunosuppressing or immunostimulatory peptide conforms to the motifs represented by the formulae of the invention, with the proviso that the peptides described in AAW3699, AAW37000 and AAW41647 to AAW41682 are excluded. An immunosuppressant can be used to treat transplant rejection or autoimmune disease, e.g. rheumatoid arthritis, systemic lupus erythematosis, Sjogren's syndrome, scleroderma, mixed connective tissue disease, dermatomyositis, polymyositis, Reiter's syndrome, Behcet's disease, type I diabetes, Hashimoto's thyroiditis,
                                                                                                                                                                                                                                                                                                                                                    Nucleic acids encoding cysteine- or methionine-containing peptide(s) which have immuno:stimulatory or immunosuppressive activity - can be to treat, e.g. cancers, infection, auto:immune disease or transplant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present peptide is a known immunomodulatory agent. It is described as a cyclic analogue of thymic humoral factor gamma-2 and was disclosed in WO9501182. The peptide is one of 6 known peptides which concord with a highly generic sequence covering immunomodulatory peptides of 4-15 amino acids being claimed in the present patent application; all 6 peptides are specifically excluded from the claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide(s) with immunomodulatory activity - useful for treating e chronic bronchitis, malignancies, herpes, AIDS or atopic disease, adjuvants in vaccines, or for inhibiting graft rejection.
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23-SEP-1996;
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                                                                                                                                                                                                                                                                                                                            rejection
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96SE-00003469
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                                                                                                                                                                                                                                                               19; 183pp; English.
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ion; autoimmune disease;
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                                                                                                                         The present invention relates to a molecule with two pendant phenylarsine moieties. The invention is useful for imparting detectable properties to target material, for detecting target material, for monitoring binding process, for isolating and immobilizing at least one target material, for screening a library of peptides and identifying a cysteine containing peptides and for labeling protein in living cells. The present sequence
                                                                                         Sequence 8 AA;
                                                                                                                                                                                                          Claim
                                                                                                                                                                                                                                           New bis-phenylarsine derivatives useful for labeling protein in living cells and for screening a library of peptides and identifying a cysteine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pendant phenylarsine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Target peptide
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                                                                                                                                                                                                                               containing peptides
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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                                                                                                                represents a target peptide of the invention.
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Pred. No. 2e+
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                                            Score 31; DB
Pred. No. 2e+
0; Mismatches
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2e+06;
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RESULT 35 ADV60194 ID ADV60 XX

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RESULT 36
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Best Local
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                                                                                                                                                                                                                                                                                              quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present sequence represents a peptide from a VEGF peptide combo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADV60194;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the family of protein.
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               ADZ28911;
                                                                                                                                                                                                                                                                   Sequence 8
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17-JUN-2003; 2003US-0479061P.
                                               ADZ28911 standard; protein;
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                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                  Score 31; DB 9;
Pred. No. 2e+06;
Mismatches
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RESULT 37
ADV60195
ID ADV60

ADV60195 standard; peptide; 9 AA

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2452

Query Match Best Local | Matches

Similarity 100 4; Conservative

100.0%; Score 31; 100.0%; Pred. No.

0;

Mismatches

0

Gaps

0

DB 9; 2e+06;

Length 8;

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The invention relates to a cellular extract from an organism comprising a CC SlyD gene, substantially free of a SlyD polypeptide that binds to a bi-CC arsenical reagent. Also described are: (i) a kit for the in vitro CC production of proteins, the kit comprising a cellular extract above; (ii) a polypeptide comprising an amino acid sequence that binds specifically to EDT 2 [4',5'.bis(1,3,2-dithioarsolan-2-yl] fluorescein-(1,2-ethanedithiol)2, preferably, the cellular extract is cequence that binds specifically to EDT 2 [5'.bis(1,3,2-dithioarsolan-2-yl) fluorescein-(1,2-ethanedithiol)2. Preferably, the cellular extract is cene. The SlyD mutant gene encodes a SlyD that is mutated in an amino acid sequence that binds a bi-arsenical molecule. The SlyD mutant gene encodes a SlyD that is mutated in an amino caid sequence that binds a bi-arsenical molecule. The SlyD mutant gene encodes a SlyD that is mutated in an amino caid sequence that binds a bi-arsenical molecule. The SlyD mutant gene encodes a slyD that is mutated in an amino caid sequence that binds a bi-arsenical molecule. The SlyD mutant gene calluced activity of at least one enzyme that catalyzes hydrolysis of high-energy phosphate bonds or hydrolysis of formation of phosphodiester bonds. It further comprises at least one inhibitor of at least one enzyme that catalyzes hydrolysis or formation of phosphodiester bonds. The cellular extract further comprises a flast one enzyme that catalyzes hydrolysis or formation of phosphodiester bonds. The cellular extract further comprises at least one inhibitor of at least one enzyme that catalyzes hydrolysis or formation protein encoding an comprises a nuclease acid encoding a fusion protein encoding an comprises a nuclease do a tag, where the tag binds a bi-arsenical comprises a nuclease of a tag, where the tag binds a bi-arsenical comprises and comprises and polypeptides. It is comprised to a tag, where the tag binds a bi-arsenical comprises and comprise and comprise and comprise and comprise and comprise and comprise a
New cellular extract from an organism comprises a SlyD gene, substantially free of a SlyD polypeptide that binds to a bi-arsenical reagent, useful for synthesizing, purifying, or detecting biomolecules,
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Sequence 8
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RESULT 38
AAR50846
ID AAR50
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AC AAR50
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                                                                                                                                                                                    Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex
   AAR50846;
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                               AAR50846 standard; peptide; 10
                                                                                                                                                                                                                                                     Sequence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-JUN-2003; 2003EP-00101775.
17-JUN-2003; 2003US-0479061P.
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4; Conserv
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Pred. No. 2e+
0; Mismatches
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                                                                                                                                                                                                    DB 9;
2e+06;
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                                                                                                                                                                                                                  Length 9;
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                                                                                                                                                                                    Gaps
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The peptide modifies the binding of P-, E- and L-selectin and of integrins, especially intercellular adhesion molecule-1 (ICAM-1). It inhibits cellular adhesion. It can be used to treat or prevent inflammation, coagulation, ischaemia, reperfusion injury, bacterial sepsis, disseminated intravascular coagulation, adult respiratory distress syndrome, tumour metastisis, rheumatoid arthritis and atherosclerosis. When labelled it can be used to detect defective selectin (or integrin) binding ligands or high (localised) concentrations of selectin (or integrin) binding cells, especially leucocytes or tumour cells. It can also be used for in vivo imaging. This is a specific example of a degenerate sequence given in the specification. (Opdated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Selectin; integrin; intercellular adhesion molecule; ICAM-1; inh inflammation; coagulation; ischaemia; reperfusion injury; bacterial sepsis; tumour metastis; disseminated intravascular coagulation; rheumatoid arthritis; adult respiratory distress syndrome; atherosclerosis; treatment; prevention; detection; tumour; leucocyte; imaging.
Sequence 10
                                                                                                                                                                                                                                       New cyclic peptide(s) inhibiting selections and integrins - preven cellular adhesion in treatment of inflammation, perfusion, injury, metastasis, etc. also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
27-SEP-1994
                                                                                                                                                                                                             Claim 2; Page 25; 175pp; English
                                                                                                                                                                                                                                                                                                                                                                                      08-SEP-1992;
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(first entry)
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Query Match

100.0%;

Score

31; DB 2;

Length 10;

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RESULT 39
AARSO868
ID AARSO
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Matches
                                                                                            New cyclic peptide(s) inhibiting selections and integrins - cellular adhesion in treatment of inflammation, perfusion, metastasis, etc. also for diagnosis.
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Misc-difference
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27-SEP-1994
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The peptide modifies the binding of P-, E- ar integrins, especially intercellular adhesion inhibits cellular adhesion. It can be used to
                                                              Claim 2; Page 26; 175pp; English.
                                                                                                                                                            WPI; 1994-100848/12.
                                                                                                                                                                                           Heavner GA;
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ches 0;
                E- and L-selectin and of esion molecule-1 (ICAM-1).
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or prevent
                                                                                                            injury,
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                                                                                                                            prevent
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RESULT 40
AAR50834
ID AAR50
XX AAR50
AC AAR50
XX 25-MA
DT 27-SE
XX Selec
XW Selec
XW
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Best Local (
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27-SEP-1994
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 WPI; 1994-100848/12
                                  Heavner GA
                                                                                               08-SEP-1992;
                                                                                                                            08-SEP-1993;
                                                                                                                                                            17-MAR-1994.
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inflammation, coagulation, ischaemia, reperfusion injury, bacterial sepsis, disseminated intravascular coagulation, adult respiratory distress syndrome, tumour metastisis, rheumatoid arthritis and atherosclerosis. When labelled it can be used to detect defective selectin (or integrin) binding ligands or high (localised) concentrations of selectin (or integrin) binding cells, especially leucocytes or tumour cells. It can also be used for in vivo imaging. This is a specific example of a degenerate sequence given in the specification. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disseminated intravascular coagulation; rheumatoid arthritis; adult respiratory distress syndrome; atherosclerosis; treatment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Selectin; integrin; intercellular adhesion molecule;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cyclic peptide inhibitor of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR50834 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prevention; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bacterial sepsis; tumour metastisis;
(CENZ ) CENTOCOR INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    coagulation; ischaemia; reperfusion injury;
                                    9208-00941653
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Pred. No. 1.9
); Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               leucocyte;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and integrins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               imaging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2
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RESULT 41
AAR50895
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The peptide modifies the binding of P-, E- and L-selectin and of integrins, especially intercellular adhesion molecule-1 (ICAM-1). It inhibits cellular adhesion. It can be used to treat or prevent inflammation, coagulation, ischaemia, repertusion injury, bacterial sepsis, disseminated intravascular coagulation, adult respiratory distress syndrome, tumour metastisis, rheumatoid arthritis and atherosclerosis. When labelled it can be used to detect defective selectin (or integrin) binding ligands or high (localised) concentrations of selectin (or integrin) binding cells, especially leucocytes or tumour cells. It can also be used for in vivo imaging. This is a specific example of a degenerate sequence given in the specification. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New cyclic peptide(s) inhibiting selections and integrins - cellular adhesion in treatment of inflammation, perfusion, i
                                                                                                             Misc-difference
                                                                                                                                     Misc-difference
                                                                                                                                                              Misc-difference
                                                                                                                                                                                      Disulfide-bond
Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
27-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        metastasis, etc. also for diagnosis.
17-MAR-1994
                      WO9405310-A1
                                                                                     Misc-difference
                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                  prevention;
                                                                                                                                                                                                                                                                                                                                            disseminated
                                                                                                                                                                                                                                                                                                                                                                                                          Cyclic peptide inhibitor of selectins and integrins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR50895 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                          bacterial sepsis; tumour metastisis;
                                                                                                                                                                                                                                                                                                                                                                                    Selectin;
                                                                                                                                                                                                                                                                                                              minated intravascular coagulation; rheumatoid arthritis; respiratory distress syndrome; atherosclerosis; treatment; ntion; detection; tumour; leucocyte; imaging.
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                                                                                                                                                                                                                                                                                                                                                                                  integrin; intercellular adhesion molecule; ICAM-1; inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ΑĄ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                     coagulation;
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                                                                                                                                                                                                                 "D-form
                                                                                                                                                                                                                                         "D-form residue."
                                                                        "D-form
                                                                                                                                                                                                                                                                                                                                                                 ischaemia; reperfusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 31;
Pred. No.
                                                                                                                                                                                                                 residue."
                                                 residue."
                                                                        residue."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 10
                                                                                                                                                                                                                                                                                                                                                                    injury;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    - prevent
injury, "
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RESULT 42
AAR50830
ID AAR50
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The peptide modifies the binding of P-, E- and L-selectin and of integrins, especially intercellular adhesion molecule-1 (ICAM-1). It inhibits cellular adhesion. It can be used to treat or prevent inflammation, coagulation, ischaemia, reperfusion injury, bacterial sepsis, disseminated intravascular coagulation, adult respiratory distress syndrome, tumour metastisis, rheumatoid arthritis and atherosclerosis. When labelled it can be used to detect defective selectin (or integrin) binding ligands or high (localised) concentrations of selectin (or integrin) binding cells, especially leucocytes or tumour cells. It can also be used for in vivo imaging. This is a specific example of a degenerate sequence given in the specification. (Updated on proper to the field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New cyclic peptide(s) inhibiting selections and integrins - cellular adhesion in treatment of inflammation, perfusion,
            Misc-difference
                                        Misc-difference
                                                                                                        Misc-difference
                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                 disseminated intravascular coagulation; rheumatoid arthritis; adult respiratory distress syndrome; atherosclerosis; treatment;
                                                                                                                                                                                                                                            inflammation; coagulation; ischaemia bacterial sepsis; tumour metastisis;
                                                                                                                                                                                                                                                                        Selectin; integrin; intercellular adhesion
                                                                                                                                                                                                                                                                                                  Cyclic peptide inhibitor of selectins and integrins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; Page 28; 175pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   metastasis, etc. also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1994-100848/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Heavner
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                                                                 Misc-difference
                                                                              Disulfide-bond
                                                                                                                                  Misc-difference
                                                                                                                                                                                                       prevention;
                                                                                                                                                                                                                                                                                                                               27-SEP-1994
                                                                                                                                                                                                                                                                                                                                             25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                       AAR50830;
                                                                                                                                                                                                                                                                                                                                                                                                  AAR50830 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2003 to correct PN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                  peptide;
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                                                                                             "D-form residue
                                                                                                                      "D-form residue.
 "D-form
                           "D-form
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            field.)
                                                                                                                                                                                                                                                       illular adhesion molecule; ICAM-1; inhibition;
ischaemia; reperfusion injury;
                                                                                                                                                                                                                                                                                                                                                                                                    10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 31; DB 2;
Pred. No. 1.9e+02;
residue
                                                      residue.
                           residue.
                                                                                                                                                                                                       leucocyte;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  perfusion, injury,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>.</u>.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               prevent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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RESULT 43
AAG94724
ID AAG94
XX AAG94
XX AAG94
XX Humar
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XX Homo
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The peptide modifies the binding of P-, E- and L-selectin and of integrins, especially intercellular adhesion molecule-1 (ICAM-1). It inhibits cellular adhesion. It can be used to treat or prevent inflammation, coagulation, ischaemia, reperfusion injury, bacterial sepsis, disseminated intravascular coagulation, adult respiratory distress syndrome, tumour metastisis, rheumatoid arthritis and atherosclerosis. When labelled it can be used to detect defective selectin (or integrin) binding ligands or high (localised) concentrations of selectin (or integrin) binding cells, especially leucocytes or tumour cells. It can also be used for in vivo imaging. This is a specific example of a degenerate sequence given in the specification. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New cyclic peptide(s) inhibiting selections and integrins - cellular adhesion in treatment of inflammation, perfusion, metastasis, etc. also for diagnosis.
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                          13-DEC-2000; 2000WO-GB004776.
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Pred. No.
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RESULT 44
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                                                                                                                                                                                                                               A set of peptide ligands consisting of specific complementary peptides proteins encoded by genes of the human genome, useful in an assay for screening and identifying of one or more novel peptides which are drug
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                                                                                                                                                                                                             or pro-drugs.
                                                                                                                                                         Page 178;
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100.0%; Pr
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Pred. No.
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The invention relates to a set of complementary peptide ligands generated from the human genome. The complementary peptides interact with their relevant target proteins encoded in the human genome. They can be used as reagents in drug discovery and as lead ligands to facilitate drug design and development. The present sequence is a complementary peptide provided

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RESULT 46
AAG86918
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AAG86916
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 10
                                                                                                                                                                                                                                                                                                                       Identifying complementary peptides by analysis of protein and nucleotide sequence databases, useful in drug design.
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                                                                                                                                                                                                                                                                                                                                                                                Roberts GW,
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                                AAG86918 standard; peptide; 10 AA.
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Pred. No. 1.9e+02;
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RESULT 47
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to the identification of complementary peptides analysis of protein and nucleotide sequence databases from higher eukaryotic genomes, excluding human and plants. The specific complementary peptides interact with their relevant target proteins encoded in the eukaryote genome. The peptides may be used as reagents drugs for drug discovery and as lead ligands for drug design and development. The present sequence is a complementary peptide from
                   17-JUN-2003; 2003EP-00101775.
17-JUN-2003; 2003US-0479061P.
                                                       17-JUN-2004; 2004WO-EP051158
                                                                                                                           Unidentified
                                                                                                                                                diagnosis; cancer; obesity; diabetes; asthma; inflammation; depression;
food; feedstuff; cosmetics; agriculture; animal breeding; VEGF.
                                                                                                                                                                                                                                 ADV60196;
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying complementary peptides by analysis of protein and nucleotide sequence databases, useful in drug design.
                                                                              23-DEC-2004.
                                                                                                                                                                                                          10-MAR-2005
                                                                                                                                                                                                                                                      ADV60196 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Saccharomyces cerevisiae
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drug discovery; drug design.
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                                                                                                    WO2004111636-A2
                                                                                                                                                                                   VEGF peptide SEQ ID NO 7701.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; Page 287; 488pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roberts GW, Heal JR;
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                                                                                                                                                                                                                                                      peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  complementary peptide; peptide identification;
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Pred. No. 1.9
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC inflammation, neuropsychiatric disorders such as depression, for CC quantifying one to several hundreds of protein disease markers CC grantifying one to several hundreds of protein disease markers CC simultaneously leading to a more accurate diagnostic sub-classification, CC for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels CC diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, CC development of improved animal models, biomarkers related with CC toxicology, clinical drug development, guidance marketed drugs, CC prognostic or diagnostic disease markers, drug target validation and CC selection, monitoring protein splicing, drug lead profiling, pathway CC analysis, answering basic disease biology questions, and in the fields of CC food and feed, cosmetics, agriculture and animal breeding. The present CC sequence represents a peptide from a VEGF peptide combo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  one or more particular proteins in a sample, for diagnosing co-
genetic diseases such as cancer, obesity, diabetes, asthma and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex
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Bergstrand H,
                                                                                                                          12-APR-1996;
23-SEP-1996;
                                                                                                                                                                                                                       04-APR-1997;
                                                                                                                                                                                                                                                                                23-OCT-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunomodulator; immunosuppressant; immunostimulator; treatment;
transplant rejection; autoimmune disease; cancer; infection.
                                                                (ASTR ) ASTRA
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   Eriksson T,
                                                                                                                          96SE-00001422
96SE-00003469
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   Lindvall M,
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Pred. No. 1.9
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1.9e+02;
0;
      Saernstrand
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding cysteine- or methionine-containing peptide(s) which have immuno:stimulatory or immunosuppressive activity - can be to treat, e.g. cancers, infection, auto:immune disease or transplant
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                                                                                                                                                                                                                                                                        Synthetic genes
                                                                                                                                                                                                                                                                                                                                                                             Bergstrand H,
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                                                                                                                                                                                               Claim 32; Page 52; 104pp;
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No.
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The present invention describes nucleic acid molecules comprising a coding sequence encoding an immunoreactive peptide and further encoding a protein targeting sequence. The nucleic acid is administered to a patient so that its expression product, an immunoactive peptide, modulates an immune response in a patient. The nucleic acid can also be used to treat cancer, either after surgery to remove a portion of the cancer or after cancer, either after surgery to remove a portion of the cancer or after

conjunction with

the nucleic acid. Cells containing the nucleic acid molecule can

also

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RESULT 50
ADV60191
ID ADV60
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         simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, promostic or diagnostic disease markers, drug target validation and promostic or diagnostic disease markers, drug target validation and
                                                                                                                                                                 The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diagnosis; cancer; obesity; diabetes; asthma; inflammation; de food; feedstuff; cosmetics; agriculture; animal breeding; VEGF
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17-JUN-2003; 2003US-0479061P.
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profiling, pathway
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CC genetic diseases such as cancer, obesity, diabetes, asthma and CC inflammation, neuropsychiatric disorders such as depression, for CC quantifying one to several hundreds of protein disease markers CC simultaneously leading to a more accurate diagnostic sub-classification, CC for determining the extent of protein modification in a particular sample CC of proteins, for tissue-typing analysis, for prenatal testing to detect CC the presence of a congenital disease or for quantitating protein levels CC diagnostic of a chromosomal abnormality, for diagnosing immune diseases CC or neurological diseases, as biomarkers preclinical drug development, CC development of improved animal models, biomarkers elated with CC toxicology, clinical drug development, guidance markere tralated with CC prognostic or diagnostic disease markers, drug target validation and CC prognostic or diagnostic disease markers, drug target validation and CC prognostic or diagnostic disease biology questions, and in the fields of cond and feed, cosmetics, agriculture and animal breeding. The present
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food and i
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stuff; cosmetics; agriculture; animal breeding; VEGF
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                     quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for pranatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present
                                                                                                                                                                                                                                                       The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
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17-JUN-2003; 2003US-0479061P
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            sequence represents a peptide
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               CC target proteins. The method comprises: (1) contacting an antibody-coated conspicient, where the fusion protein, where the antibodies bind specifically to the fusion protein and the fusion protein comprises at the fusion protein protein protein comprises at the class a portion of the target protein linked to a carrier protein, and comprises at the presence of the presence of the carrier protein, and comprises at the protein, where the presence of a carrier protein indicates the presence of an antibody to the target protein. The target protein may be an isoform of a protein that is associated with disease, e.g. Vascular candidates associated with cancer; prostate Specific Antigen (RADW28645), which are associated with prostate cancer and WEGF121 (ADW28645), which is associated with prostate cancer and Her2 (MADW28644), which is associated with prostate cancer. Also claimed are methods for: generating (M2) monoclonal antibodies, where each monoclonal comprises isolating (M3) an antibody binding specifically at least one monoclonal comproses in from several antibody binding specifically at least one monoclonal comproses in from several antibody; determining (M4) the presence of an antibody ending specifically to a target protein from several antibody; determining (M3) an antibody to a particular molecule is consisted which the presence of an antibody to a particular molecule is consisted with the presence of an antibody and antibody to a particular molecule is consisted with the presence of an antibody and antibody and antibody to a particular molecule is consisted with the presence of an antibody and antibody as a particular molecule is consisted with the presence of an antibody and antibody to a particular molecule is consisted with the presence of an antibody and antibody to a particular molecule is consisted with the presence of an antibody and antibody to a particular molecule is consisted with the presence of an antibody and antibody to a particular molecule is consisted with antiparticular molecule is con
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying antibody to protein, involves contacting solid surface coated with antibodies with fusion protein having portion of target protein and carrier protein and conducting assay to determine presence of carrier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; Antirheumatic; Antidiabetic; Cardiant; Nootropic; Neuroprotective; antibody identification; antibody production; antibody therapy; cancer; vascular endothelial cell growth factor; VEGF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to a method for identifying antibodies for target proteins. The method comprises: (i) contacting an antibody-coated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 32; SEQ ID NO 2; 55pp; English.
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to particular cells e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
  cancer cell.
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Best Local S
Matches 4
The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants of one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obseity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    diagnosis; cancer; obesity; diabetes; asthma; inflammation; depression; food; feedstuff; cosmetics; agriculture; animal breeding; GPCR.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-JUN-2004; 2004WO-EP051158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               agriculture, comprises generating peptides family of protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2005-057893/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kas K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-JUN-2003; 2003EP-00101775
17-JUN-2003; 2003US-0479061P
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                                                                                                                                                                                                                                                                                                                                                                                                                           Example; SEQ ID NO 2373; 265pp; English.
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Pred. No.
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            The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants of one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present sequence represents a peptide from a G-protein coupled receptor peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G protein coupled receptor peptide SEQ ID NO 2489.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-JUN-2003; 2003EP-00101775
17-JUN-2003; 2003US-0479061P
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                                                                                                                                                                                                                                         Example; SEQ
                                                                                                                                                                                                                                                                                         agriculture, comprises generating peptides by applying a
                                                                                                                                                                                                                                                                                                       Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in
                                                                                                                                                                                                                                                                                                                                                            WPI; 2005-057893/06
                                                                                                                                                                                                                                                                                                                                                                                            Kas K,
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Pred. No. 2.8e+02;
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RESULT 56
ADV55097

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Best Local S
Matches 4
The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants of one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for quantifying one to several hundreds of protein diseases markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying a peptide of useful for diagnosing a agriculture, comprises
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17-JUN-2003;
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food; feedstuff; co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example;
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2003US-0479061P.
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cosmetics; agriculture; anima
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a variety of diseases, drug development or in
s generating peptides by applying a digest on the
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cardiovascular disease; angina pectoris; arteriosclerosis; heart failure; brain disorder; brain abscess; meningitis; Alzheimer's disease; sarcoma; cytostatic; carcinoma; cardiant; neuroprotective; antiinflammatory;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    head; neck;
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The invention relates to human thioredoxin nucleic acid and polypeptide molecules, designated 22108 and 47916. The compound that modulates the activity or expression of 22108 and 47916 nucleic acid is useful for treating or preventing a disorder characterised by aberrant activity of 22108 and 47916-expressing cell, specifically for reducing or inhibiting the aberrant activity of the 22108 and 47916 expressing cancer cell. The 22108 and 47916 nucleic acid and polypeptide are useful for diagnosing, preventing or treating cancer in a subject (e.g. carcinoma, sarcoma, metastatic or haematopoietic disorders (e.g.leukaemia), or cancers of the lung, breast, thyroid, head neck, prostate or genito-urinary tract), cardiovascular diseases (e.g. angina pectoris, arteriosclerosis or heart failure) or brain disorders (e.g. brain abscess, meningitis, Alzheimer's diseases). The thioredoxin DNA is also useful in gene therapy. The

New human thioredoxin nucleic acid and polypeptide molecules, or 22108 and 47916, useful for diagnosing, preventing or treating (e.g. carcinoma), cardiovascular diseases (e.g. heart failure)

designated or bra

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124pp; English

WPI; Bandaru R, (MILL-) MILLENIUM PHARM INC

Kapeller-Libermann

25-SEP-2000;

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RESULT 58
ADV60188
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       quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present sequence represents a peptide from a VEGF peptide combo.
                                                                                                                                                                                                                                                     genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
                                                                                                                                                                                                                                                                                   corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex
                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the family of protein.
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                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of identifying a peptide combo which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2005-057893/06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vandekerckhove J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CGPC 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         peptide; 20 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NO 7693
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Krols L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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Best Local (
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                         quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for granatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic ofa chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as blomarkers preclinical drug development, development of improved animal models, blomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present sequence represents a peptide from a VEGF peptide combo.
                                                                                                                                                                                                                                                                             The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants of one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2005-057893/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kas K,
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17-JUN-2003; 2003US-0479061P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-JUN-2004; 2004WO-EP051158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VEGF peptide SEQ ID NO 7694.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-MAR-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADV60189;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-DEC-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis; cancer; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        family of protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (VIBV-) VIB VZW.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          feedstuff; cosmetics;
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                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 7694; 265pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CGPC 17
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100.0%; Pred. No. 3.2e+02;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Krols L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diabetes; asthma; inflammation; depression; agriculture; animal breeding; VEGF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
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Sequence 21 AA

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DB 9;

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RESULT 60
AD338160
ID AD338160
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                                               invention. Human secreted proteins have activities based on the tissues and cells the genes are expressed in Example of activities include: immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; and ophthalmological. The polymucleotides and polymeptides can be are used to grevent, treat or ameliorate a medical condition in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. They are also used in diagnosing a pathological condition or susceptibility to a pathological condition or susceptibility to a pathological condition or susceptibility to a disorders, respective and pathological condition or susceptibility of a disorders, infections caused by bacteria, viruses and fungi and ocular disorders. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to sunburn, to maintain organs before transplantation, for supporting cell culture of the polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 4; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; ophthalmological; gene therapy; autoimmune disease; infection; hyperproliferative disorder; cardiovascular disorder; anglogenesis; cerebrovascular disorder; nervous system disorder; ocular disorder; wound healing; skin aging; food additive; preservative.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The polynucleotide sequences given in AAC69399 to AAC69445 encode human secreted proteins given in AAB38119 to AAB38165. AAB38166 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acids encoding human secreted proteins, used to prevent, ameliorate, or diagnose conditions such as autoimmune disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB38160
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB38201 represent human secreted polypeptide sequences and proteins homologous to them, which are given in the exemplification of the pro-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 11; Page 358; 374pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rosen CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ruben SM,
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99US-0171550P
   also
                                  to regenerate tissues and in chemotaxis.
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Pred. No.
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   preservative to
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quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosting immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present
                                                                                                                                                                                                             The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
                                                                                                                                                                                                                                                                                                                                                                                               Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    increase or decrease storage capabilities. AAC69390 to AAC69398 and AAB38118 represent sequences used in the exemplification of the present
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17-JUN-2003; 2003US-0479061P
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Pred. No.
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                simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present
                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
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17-JUN-2003; 2003US-0479061P.
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                                                                                                                                                                                                                                                                                                                                                                                                quantifying one to several hundreds of protein disease markers
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from a G-protein coupled receptor
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            of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels disgnostic of a chromosomal abnormality, for disgnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or disgnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present sequence represents a peptide from a G-protein coupled receptor peptide
                                                                                                                                                                                                                                                        The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G protein coupled receptor peptide SEQ
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17-JUN-2003; 2003US-0479061P
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Pred. No. 3.6e+02;
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                                        The invention relates to a method of identifying a peptide combo which CC corresponds with a family of proteins where each of the members of the CC peptide combo is derived from a unique protein from the family. The CC one or more particular proteins in a sample, for diagnosing complex CC genetic diseases such as cancer, obesity, diabetes, asthma and CC inflammation, neuropsychiatric disorders such as depression, for CC quantifying one to several hundreds of protein disease markers CC simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample CC of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels CC diagnostic of a chromosomal abnormality, for diagnosing immune diseases CC or neurological diseases, as biomarkers preclinical drug development, CC development of improved animal models, biomarkers related with CC concentration of the protein splicing, drug target validation and CC geneticn, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADV54929
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G protein coupled receptor peptide SEQ ID NO 2426
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-JUN-2003; 2003EP-00101775.
17-JUN-2003; 2003US-0479061P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diagnosis; cancer; obesity; diabetes; asthma; inflammation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       feedstuff;
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4; Conserv
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Pred. No.
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RESULT 65
ADV55148
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CC genetic diseases such as cancer, obesity, diabetes, asthma and cc inflammation, neuropsychiatric disorders such as depression, for cquantifying one to several hundreds of protein disease markers cc simultaneously leading to a more accurate diagnostic sub-classification, for cf for determining the extent of protein modification in a particular sample composition for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels cc diagnostic of a chromosomal abnormality, for diagnosing immune diseases cor neurological diseases, as biomarkers preclinical drug development, cc development of improved animal models, biomarkers related with coxicology, clinical drug development, guidance marketed drugs, compositic or diagnostic disease biology questions, and in the fields of cod and feed, cosmetics, agriculture and animal breeding. The present composition of the present application coupled receptor peptide from a G-protein coupled receptor peptide.
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                                                                                                                                                                                                                                                                                                                           corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G protein coupled
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17-JUN-2003; 2003US-0479061P
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Example; SEQ ID NO 2645; 265pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   family of protein.
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RESULT 66
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                                                                                                                            Sequences AAB37984-B38019 represent the amino acid sequences of 27 human composed proteins encoded by the genes AAC69084-C69119. The genes and conditions, e.g. by protein or gene therapy. The genes are isolated from caring of human tissues disclosed in the specification. The nucleic cardis, proteins, antibodies and (ant)agonists are useful in the cardis, proteins, antibodies and (ant)agonists are useful in the cardis, proteins, antibodies and (ant)agonists are useful in the covarian cancer, and other cancers of the adrenal gland, bone, bone immune disorders e.g. Addison's disease, altergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f) infectious diseases such as viral, bacterial, fungal and parasitic
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Best Local Similarity
Matches 4; Conserv
                               Matches
                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acid molecules encoding 27 human secreted proteins for diagnosing, preventing, treating or ameliorating medical conditions and used as food additives or preservatives.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antiulcer; vulnerary; antionvulsant; antibacterial; antifugal; antiparastic; cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder;
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                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                           Disclosure;
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                                             Local Similarity
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Lafleur DW,
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Moore PA,
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Pred. No. 3.8e+02;
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Pred. No. 3.6e+02;
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Olsen
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1 HS, Young
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PE;
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Matches Query Match Best Local

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                                                                                        quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a congenital disease or for quantitating protein levels or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, aprognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present sequence represents a peptide from a G-protein coupled receptor peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            genetic diseases such as cancer, obesity, diabetes, asthma an inflammation, neuropsychiatric disorders such as depression,
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Sequence
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17-JUN-2003; 2003US-0479061P.
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(UYGE-) UNIV GENT
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                                                                                                                                                                                                      The invention relates to a method of identifying a peptide combo which CC corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The CC peptide combo is useful for quantifying specific known splice variants of CC one or more particular proteins in a sample, for diagnosing complex CC genetic diseases such as cancer, obesity, diabetes, asthma and CC inflammation, neuropsychiatric disorders such as depression, for CC quantifying one to several hundreds of protein disease markers CC genetic sub-classification, for cc simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample CC of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels CC diagnostic of a chromosomal abnormality, for diagnosing immune diseases CC or neurological diseases, as biomarkers preclinical drug development, CC development of improved animal models, biomarkers related with CC toxicology, clinical drug development, guidance marketed drugs, answering basic disease markers, drug target validation and ce selection, monitoring protein splicing, drug target validation and feed, cosmetics, agriculture and animal breeding. The present
                                                                                    Query Match
Best Local Similarity
                                                                    Matches
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17-JUN-2003; 2003US-0479061P
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food; feed:
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                                                                                                                                                                           sequence represents a peptide combo.
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CGPC
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cosmetics; agriculture; animal breeding; GPCR.
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Pred. No. 3.8e+02;
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CGPC 11

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Matches Query Match Best Local |

4. Similarity

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Mismatches

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Gaps

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100.0%;

Score 31; DB 9; Pred. No. 3.8e+02;

Length 25; Indels

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RESULT 69
ADV54928
The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is derived from a unique protein from the family. The competition of the genetic diseases such as cancer, obesity, diabetes, asthma and competition, neuropsychiatric disorders such as despression, for cumulifying one to several hundreds of protein disease markers and complex simultaneously leading to a more accurate diagnostic sub-classification, for complex complex computation, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnostic immune diseases or neurological diseases, as biomarkers preclinical drug development, component of improved animal models, biomarkers related with coxicology, clinical drug development, guidance markered drugs, analysis, answering protein splicing, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway componed represents, agriculture and animal breeding. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
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feedstuff; cosmetics; agriculture; animal breeding; GPCR.
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RESULT 70
ADC89744
ID ADC89
                                                                                                            The invention relates to a cyclic VEGF (vascular endothelial growth Cf factor) peptide (ADC89740) with NP-1 (neuropilin-1) antagonist activity. Cf factor) peptide corresponds to residues 138-165 of the VEGF-165 splice variant, but contains 2 disulphide bonds. It acts as a competitive cinhibitor of NP-1, which plays an important role in angiogenesis (including VEGF-mediated tumour angiogenesis), neurodegeneration Cf (including VEGF-mediated tumour angiogenesis), neurodegeneration NP-1 is a non-tyrosine kinase receptor for the VEGF-165 splice variant. Cf Additionally, VEGF is known to promote the survival of tumour cells that express NP-1, including some breast carcinoma cells. The cyclic VEGF cepptide of the invention can be used to stimulate nerve growth and Cf repair, to treat neurodegeneration, and for anti-cancer therapy. It can also be used for treating other angiogenic conditions such as eye conceases and rheumatoid arthritis, and as an immunosuppressant for use in transplantation. The present sequence represents a cyclic VEGF-165 concerns the conditions of VEGF-165 to NP-1, unlike the cyclic peptide of the invention conditions conditio
Query Match 100.0%; Score 31; DB 7; Best Local Similarity 100.0%; Pred. No. 4.1e+02; Matches 4; Conservative 0; Mismatches 0;
                                                                                                      Sequence 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 1; 21pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New cyclic peptide derived from vascular endothelial growth factor, useful for treating e.g. neurodegeneration and cancer, antagonizes neuropilin-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Selwood D, Loehr M, Zachary I;
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Unidentified.
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Disulfide-bond
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submitted to the Protein Sequence Database, October 200 A; Description: Purification and Sequence determination A; Reference number: A59356 A; Accession: A59356 A; Accession: A59356 A; Molecule type: protein
                                                                                                       neurotoxin BmK 41-2 - scorpion (Buthus Martensii Karsch) C;Species: Buthus Martensii Karsch C;Date: 31-Dec-2001 #sequence_revision 31-Dec-2001 #text C;Accession: A59356 R;Xu, C.
                                                                                                                                                    RESULT
A59356
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C;Superfamily: scorpion neurotoxin
C;Keywords: neurotoxin; venom
F;1-18,4-25,15-30,19-32/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                               A;Accession: A01758
A;Molecule type: protein
A;Residues: 1-35 <ROS>
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                                                                                                                                                                                                                                                                                                                                                                                       neurotoxin P2 -
                                           Status: preliminary
Molecule type: protein
Residues: 1-35 <XUC>
;1-35/Product: neurotoxin BmK 41-2 #status exper;1-18,4-25,15-30,19-32/Disulfide bonds: #status
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A;Cross-references: UNIPROT:Q01644; UNIPARC:UPI00001246FE; EMBL:X67703; A;Note: the authors translated the codon TGC for residue 55 as Thr A;Note: sequence extracted from NCBI backbone (NCBIN:74217, NCBIP:74222) C;GenetLts: A;Gene: Mst84Dc
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C; Superfamily: fruit fly
C; Keywords: spermatogenes
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A;Title: A cluster of four genes selectively expressed in A;Reference number: A56565; MUID:92102953; PMID:1684716
A;Accession: S25774
                                                                                     A;Cross-references: FlyBase:FBgn0002862
C;Superfamily: fruit fly testis-specific
C;Keywords: sex-specific protein; testis
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C;Species: Drosophila melanogaster
C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 09-Jul-2004
C;Accession: S25774; C56565
                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-56 < KUH >
                                                                                                                                                                                                              A; Reference number: A; Accession: S00340
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A; Residues: 1-55 < KUH>
                                                                                                                                      A;Gene: FlyBase:Mst87F
                                                                                                                                                  C; Genetics:
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Conservative 0; Mismatches
                             Conservative
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Pred. No. 81;
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                                           Score 31;
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A;Cross-references: UNIPARC:UPI00000C4E89; GB:AL162755; GB:AL157959; NID:g7379742; PIDN A;Experimental source: serogroup A, strain Z2491
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Title: Complete DNA sequence of a serogroup A strain of Neisseria menigitidis Z2491.
A;Reference number: A81775; MUID:20222556; PMID:10761919
A;Accession: B81888
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Science 287, 1809-1815, 2000
A;Authors: Grandi, G.; Sun, L.; Smith, H.O.; Fraser, C.M.; Moxon, E.R.; Rappuoli, F. A;Title: Complete genome sequence of Neisseria meningitidis serogroup B strain MC55 A;Reference number: A81000; MUID:20175755; PMID:10710307
A;Accession: H81133
                      RESULT
S25772
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A;Title: Partial cDNA sequence for the donkey chorionic A;Reference number: S15676; MUID:90262634; PMID:2344391
A;Accession: S15676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Keywords: iron; metalloprotein F;3-49/Domain: rubredoxin homolog F;6,9,39,42/Binding site: iron (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: UNIPROT:Q9JQN3; UNIPARC:UPI00000C4E89; GB:AE002450; GB:AE002098; NII A;Experimental source: serogroup B, strain MC58 R;Parkhill, J.; Achtman, M.; James, K.D.; Bentley, S.D.; Churcher, C.; Klee, S.R.; Morel; Holroyd, S.; Jagels, K.; Leather, S.; Moule, S.; Mungall, K.; Quail, M.A.; Rajandream, Nature 404, 502-506, 2000
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A; Residues: 1-62 <LEI>
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R;Leigh, S.E.A.; Ste
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C;Superfamily: rubredoxi
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A; Residues: 1-56 < PAR>
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A; Residues: 1-56 <TET>
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testis-specific protein Mst84Da -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Species: Equus asinus (donkey)
;Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 09-Jul-2004;
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Best Local (
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fruit fly (Drosophila melanogaster)
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P 4
A;Accession: A60136
A;Molecule type: mRN
A;Residues: 1-73 <WJ
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A; Residues: 1-63 < KUH>
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A; Residues: 1-68 < KUH>
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C;Species: Drosophila melanogaster
C;Date: 26-Uul-1996 #sequence revision 26-Uul-1996 #text_change 09-Jul-2004
C;Accession: S25772; A56565
R;Kuhn, R.; Kuhn, C.; Boersch, D.; Glaetzer, K.H.; Schaefer, U.; Schaefer, M.
Mech. Dev. 35, 143-151, 1991
A;Title: A cluster of four genes selectively expressed in the male germ line of Drosoph
A;Reference number: A56565; MUID:92102953; PMID:1684716
A;Accession: S25772
Biochim. Biophys. Acta 824, 201-208, 1985
A;Title: Isolation and characterisation of keratin mRNA
A;Reference number: A60136; MUID:85122780; PMID:2578818
                                                                                                                               keratin, scale (clone CSK9) - chicken (fragment) C;Species: Gallus gallus (chicken) C;Date: 30-Jan-1993 #sequence_revision 12-Mar-19 C;Accession: A60136
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C; Superfamily: fruit fly testis-specific protein
C; Keywords: spermatogenesis; tandem repeat
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A;Title: A cluster of four genes selectively expressed in the male germ line of Drosoph' A;Reference number: A56565; MUID:92102953; PMID:1684716
A;Accession: S25775
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                                                                                          R; Wilton, S.D.; Crocker, L.A.; Rogers,
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extracted from NCBI backbone (NCBIN:74219,
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3-151, 1991
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    fruit fly (Drosophila melanogaster)

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A;Title: A cluster of four genes selectively expressed in the male germ line of A;Reference number: A56565; MUID:92102953; PMID:1684716
A;Accession: S25773
A;Molecule type: DNA
A;Residues: 1-74 KUH-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:Q9YH54
C;Superfamily: feather keratin
C;Keywords: tandem repeat
F;1-49/Region: 13-residue repeats
                                                                                                                                                                                                                                                                        R;Zhang, F.P.; Rannikko, A.; Huhtaniemi, I.
Biochem. Biophys. Res. Commun. 210, 858-865, 1995
A;Title: Isolation and characterization of testis-speci.
A;Reference number: I52320; MUID:95283549; PMID:7763258
A;Accession: I65235
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165235
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A;Map position: 3
C;Superfamily: fruit fly testis-specific protein
C;Keywords: spermatogenesis; tandem repeat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          testis-specific protein Mst84Db - fruit fly (Drosophila melanogaster) C;Species: Drosophila melanogaster C;Species: Drosophila melanogaster C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 09-JuC.hCcasein. S2873: BEEEEE
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                                                                                                                                                                                                                                      A;Status: preliminary; translated A;Molecule type: mRNA
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C; Genetics:
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                                                                                                                                                                                                                          A; Residues: 1-80 < RES>
                                                                                                                                                                                                                                                                                                                                                                     ;Species: Rattus norvegicus (Norway ra;Date: 29-May-1998 #sequence_revision
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                                                                                                          Query Match
Best Local
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Pred. No. 1.1e+02;
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Pred. No.
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A;Experimental source: calamus
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thioredoxin 2 - slime mold (Dictyostelium discoideum) (fragment)
C;Species: Dictyostelium discoideum
C;Species: Dictyostelium discoideum
C;Date: 28-Oct-1994 #sequence_revision 28-Oct-1994 #text_change 05-Oct-2004
C;Accession: B46264
R;Metterauer, B.; Jacquot, J.P.; Veron, M.
J. Biol. Chem. 267, 9895-9904, 1992
A;Title: Thioredoxins from Dictyostelium discoideum are a developmentally re A;Reference number: A46264; MUID:92250653; PMID:1577820
A;Accession: B46264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RING, W.V.; Actions, Freitas, T.
Jung, K.H.; Alam, M.; Freitas, T.
Jung, K.H.; Alam, M.; Freitas, T.
Proc. Natl. Acad. Sci. U.S.A. 97, 12176-12181, 2000
A;Authors: Hou, S.; Daniels, C.J.; Dennis, P.P.; Omer, A.D.;
A;Title: Genome sequence of Halobacterium species NRC-1.
A;Title: Genome sequence of MUID:20504483; PMID:11016950
                                                                                                            C;Accession: A02850
R;O'Donnell, I.J.; Inglis, A.S.
Aust. J. Biol. Sci. 27, 369-382, 1974
A;Title: Amino acid sequence of a feather keratin from A;Reference number: A02850; MUID:75054009; PMID:4429491
                                                                                                                                                                                                                             keratin, feather - silver gull
keratin, feather - silver gull
C;Species: Larus novaehollandiae (silver gull)
C;Species: ^ ~~~~1984 #sequence_revision 24-Apr-1984
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C;Superfamily: thioredoxin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Species: Halobacterium sp. NRC-1
C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 05-Oct-2004
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A; Residues: 1-88 < WET>
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                        A; Molecule type: protein
A; Residues: 1-98 < ODO >
A; Cross-references: UNIPE
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A; Residues: 1-89 <STO>
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                                                                                             A;Accession: A02850
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UNIPROT: P02451; UNIPARC: UPI000012E06F
ce: calamus of wing feathers from a si
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Pred. No. 1.2
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a single bird

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R;Shirai, M.; Hirakawa, H.; Kimoto, M.; Tabuchi, M.; Kishi, F.; Ouchi, K.; Shiba, Nucleic Acids Res. 28, 2311-2314, 2000
A;Title: Comparison of whole genome sequences of chlamydia pneumoniae J138.
A;Reference number: A86491; MUID:20330349; PMID:10871362
A;Accession: H86572
                                                                                                                                                                                                                                                                                                        thioredoxin [imported] - Chlamydophila pneumoniae (strain J138)
C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
C;Accession: H86572
                                                                                                                                                                                                                                                                                                                                                                                        RESULT
H86572
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                                                               A;Gene: trxA
C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                              A;Cross-references: UNIPROT:Q9Z7P5; UNIPARC:UPI000004461A; A;Experimental source: strain J138
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A; Experimental source: strain
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Science 270, 397-403, 1995
A;Title: The minimal gene complement of Mycoplasma genitalium.
A;Reference number: A64200; MUID:96026346; PMID:7569993
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A; Residues: 1-102 <STO>
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Keywords: redox-active disulfide;
;S-90/Domain: thioredoxin homology <THR>
;3-93/Disulfide bonds: redox-active #status pr
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                 100.0%;
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C;Species: Mycoplasma pneumoniae
A;Variety: ATCC 29342
C;Date: 27-Feb-1997 #segments
C;Accession
thioredoxin CP0088 [imported] - Chlamydophila pneumoniae (strains CWL029 and C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 05-Oct-2004 C;Accession: D72052; E81614
                                                                                                                                                                                                                                                                                                                                 A;Gene: trxA
C;Superfamily: thioredoxin; thioredoxin homology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Himmelreich, R.; Hilbert, H.; Plagens, H.; Pirkl, E.; Li, B.C.; Herrmann, R. Nucleic Acids Res. 24, 4420-4449, 1996
A;Title: Complete sequence analysis of the genome of the bacterium Mycoplasma A;Reference number: S73327; MUID:97105885; PMID:8948633
A;Accession: S73896
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C;Superfamily: thioredoxin; thioredoxin homology
F;9-90/Domain: thioredoxin homology <TXN>
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A; Residues: 1-102 < ARN>
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70; MUID:99000809; PMID:978413
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Pred. No. 1.3e+02;
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thioredoxin II - yeast (Saccharomyces cere N;Alternate names: protein L1933; protein C;Species: Saccharomyces cerevisiae C;Date: 31-Dec-1991 #sequence revision 31-15;Accession: S15048; A39847. Sieron 2015
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C;Superf
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A;Cross-references: UNIPROT:Q9Z7P5; UNIPARC:UPI000004461A; GB:AE001648; GB:AE001363; NII
A;Cross-references: Strain CWL029
A;Experimental source: strain CWL029
R;Read, T.D.; Brunham, R.C.; Shen, C.; Gill, S.R.; Heidelberg, J.F.; White, O.; Hickey, C.; Dodson, R.; Gwinn, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McClarty, G.; Salzberg, Nucleic Acids Res. 28, 1397-1406, 2000
A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39.
A;Reference number: A81500; MUID:20150255; PMID:10684935
A;Reference E81614
A;Status: preliminary
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C:Genetics:
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C;Species: Chlamydia muridarum, Chlamydia trachomatis MoPn
C;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 05-Oct-2004
C;Accession: C81660
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A;Accession: D72052
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R;Gan,
J. Biol
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A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39.
A;Reference number: A81500; MUID:20150255; PMID:10684935
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A; Residues: 1-102 < REA>
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Superfamily: thioredoxin; thioredoxin homology
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  Biol.
                           Z.R.
                                                                                                                                                                                             21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TC0826
  Chem.
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  266, 1692-1696,
                                                                                                                                             yeast (Saccharomyces cerevisiae)
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Pred. No. 1.3e+02;
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Pred. No. 1.36
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                                                 S64870; B38669
                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                          31-Dec-1991
                                                                                                                       YLR043c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.3e+02;
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R; Koetter, P.; Rose, M.; Entlan, K.D.
submitted to the Protein Sequence Database,
A; Reference number: S64863
A; Accession: S64870
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT
T39085
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                                                                                                                                                                                                                                                                                                                                                                                                           N;Alternate names: thioredoxine 2
C;Speciles: Schizosaccharomyces pombe
C;Date: 03-Dec:1999 #sequence revision 03-Dec-1999 #text_change 09-Jul-2004
C;Accession: T39085; T46567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Keywords: redox-active disulfide F;2-103/Product: thioredoxin II #status predicted F;9-90/Domain: thioredoxin homology <THR>
                                                                                                               A;Experimental source: strain 972h-; cosmid c7D4
R;Lenaers, G.; Perret, E.; Bonin, O.; Picard, A.; Caput, submitted to the EMBL Data Library, October 1997
A;Description: TRX2, a fission yeast stress protein.
                                                                                                                                                                                                                                                                                                                                                            R;Gentles, S.; Churcher, C.M.; Wood, V.; Barrell, submitted to the EMBL Data Library, September 199
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A;Cross-references: SGD:S0004033; MIPS:YLR043c
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A; Residues: 1-103 < KOE>
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                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-103 <GEN>
                                                                                                                                                                                                                                                                                      A;Status: preliminary; translated from GB/EMBL/DDB.
                                                                                                                                                                                                                                                                                                               A; Accession: T39085
                                                                                                                                                                                                                                                                                                                                     A; Reference number: Z21826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thioredoxin II -
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                           A; Molecule type: mRNA
                                               A; Status: preliminary; translated
                                                                          A; Accession:
                                                                                              A; Reference number: Z23075
                                                                                                                                                                                                                   A;Cross-references: UNIPROT:O14463; UNIPARC:UPI000016205F; EMBL:Z99532; PIDN:CAB16724.1
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                                                      from GB/EMBL/DDBJ
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Pred. No. 1.3e+02;
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A;Title: Yeast thioredoxin genes.
A;Reference number: A38669; MUID:91107668; PMID:1988444
A;Reference number: A38669;
A;Accession: S15048
A;Molecule type: DNA
A;Residues: 1-103 <GAN>
A;Cross-references: UNIPROT:P22217; UNIPARC:UPI000004F943; EMBL:M59169; NID:g173027; PII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GB:M62647; NID:g173047; PIDN:AAA35177.1; Reichard, P.
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R;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.; Baquero, F.; Dominguez-Bernal, G.; Duchaud, E.; Durand, L.; Dussurget, O.; E D.; Jones, L.M.; Karst, U. Science 294, 849-852, 2001
Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, A;Authors: Kreft, J.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J. A;Title: Comparative genomics of Listeria species, A;Reference number: AB1077; MUID:21537279; PMID:11679669
A;Accession: AC1582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    D.; Joneš, L.M.; Karst, U. Science 294, 849-852, 2001
Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuln, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; A;Authors: Kreft, J.; Kiln, M.; Tierrez, A.; Vazquez-Boland, J.A.; ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; A;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
                                                                                                                                                                                                                                                                                      C;Species: Listeria innocua
C;Date: 27-Nov-2001 #sequence_revision
C;Accession: AC1582
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C;Superfamily:
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A;Molecule type: DNA
A;Residues: 1-103 <GLA>
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Pred. No. 1.3e+02;
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Pred. No. 1.3e+02;
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Voss, H.; Wehland
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Voss, H.; We
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Fsihi, H.
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RESULT 27
B37192
thioredoxin - Bacillus subtilis
C;Species: Bacillus subtilis
C;Species: Bacillus subtilis
C;Date: 31-Jan-1992 #sequence revision 26
C;Accession: B37192; H69726
R;Chen, N.Y.; Zhang, J.J.; Paulus, H.
J. Gen. Microbiol. 135, 2931-2940, 1989
                                                                                                                                                                                                                                                                                                                                                                                             R.Johnson, T.C.; Yee, B.C.; Carlson, D.E.; Buchanan, B.B.; Johnson, R.S.; Mathews, J. Bacteriol. 170, 2406-2408, 1988
A;Title: Thioredoxin from Rhodospirillum rubrum: primary structure and relation to A;Reference number: A28215; MUID:88198045; PMID:3129411
A;Accession: A28215
A;Molecule type: protein
A;Residues: 1-104 <JOH-
A;Residues: 1-104 <JOH-
A;Cross-references: UNIPROT:P10473; UNIPARC:UPI000171E24
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
E;7-90/Domain: thioredoxin homology <THR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Rhodospirillum rubrum
C;Date: 28-Aug-1999 #sequence revision 26-May-1994 #text_change 05-Oct-2004
C;Accession: A28215
C;Accession: A28215
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A;Molecule type: protein A;Residues: 1-104 <JOH>
A;Residues: 1104 <JOH>
A;Cross-references: UNIFROT:P08628; UNIFARC:UPI0000136E96 C;Superfamily: thioredoxin; thioredoxin homology C;Keywords: redox-active disulfide F;8-91/Domain: thioredoxin homology <THR>
F;31-34/Disulfide bonds: redox-active #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 30-Jun-1989 #sequence_revision 26-May-1994
                                                                                                                                                                                                                                                                                                                                                                              F;29-32/Disulfide bonds: redox-active
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hem. 263, 9589-9597, 1988
mino acid sequence of thioredoxin
e number: A28086; MUID:88257078; F
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Pred. No. 1.3
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Pred. No. 1.3e+02;
; Mismatches 0;
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A;Cross-references: UNIPROT:P14949; UNIPARC:UPI000006087B; GB:U03294; GB:M26384; NID:g14
R;Kunst, F., Ogasawara, N.; Moszer, I., Albertini, A.M.; Alloni, G.; Azevedo, V.; Berter
C.; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Chc
A; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E.
Nature 390, 249-256, 1997
A;Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Galler
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Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois,
A;Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Mauceel
y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetelle
Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon,
A;Authors: Schleich, S.; Schroceter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Serot
akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Uchiyama,
T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, K.
A;Authors: Yoshikawa, H.F.; Zumstein, E.; Yoshikawa, H.; Danchin, A.
A;Title: The complete genome sequence of the Gram-postive bacterium Bacillus subtilis.
A;Reference number: A69580; MUID:98044033; PMID:9384377
A;Accession: H69726
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A;Cross-references: UNIPARC:UPI000006087B; GB:Z99118; GB:AL009126; NID:g2635200; PIDN:CA
A;Experimental source: strain 168
C;Genetics:
Eur. J. Biochem. 23, 328-335, 1971
A;Title: Yeast thioredoxin. Amino-acid sequence
A;Reference number: S05793; MUID:72100583; PMID:
                                                                 A;Residues: 1-104 <MUL>
A;Cross-references: UNIPARC:UPI000004F91C;
R;Hall, D. E. Baldesten, A.; Holmgren, A.;
Eur. J. Biochem. 23, 328-335, 1971
                                                                                                                                                                                                                           A; Reference number: A39847;
A; Accession: B39847
                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: UNIPROT:P22803; UNIPARC:UPI000004F91C; R;Muller, B.G.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             J. Biol. Chem. 266, 1692-1696, 1991
A;Title: Yeast thioredoxin genes.
A;Reference number: A38669; MUID:91107668; PMID:1988444
A;Accession: S15049
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A; Map position:
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                                                                                                                                                                                          Molecule type: DNA
                                                                                                                                                                                                                                                        . Biol. Chem. 266, 9194-9202, 1991;Title: Thioredoxin deficiency in yeast prolongs S phase and Reference number: A39847; MUID:91225027; PMID:2026619
                                                                                                                                                                                                                                                                                                                                                                                                                  ;Molecule type: DNA
;Residues: 1-104 <GAN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 05-Oct-2004;
Accession: S15049; B39847; S05793; S53932; S61947; S64531; S63858; A38669;
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; 9-90/Domain: thioredoxin homology <THR>
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Residues: 1-104 <C
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Pred. No. 1.3e+02;
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                                                                                                  GB:M62648; NID:g173049; Reichard, P.
             PMID: 4945270
                                       around the active-center disulfide
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A; Residues: 1-104 < GUW>
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A; Residues: 1-104 < GUE>
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A; Residues: 1-104 < GUF>
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4:
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A;Cross-references: UNIPARC:UPI000004F91C; EMBL:Z49133; NID:g790489; A;Experimental source: strain S288C R;Song, J.M.; Cheung, E.; Rabinowitz, J.C. submitted to the EMBL Data Library, November 1995 A;Description: Analysis of the 15.6-kb fragment encompassing the ADE3 A;Reference number: S61947 A;Accession: S61947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: protein
A;Residues: 2;27-43;98-104 <HAL>
A;Residues: 2;27-43;98-104 <HAL>
A;Residues: 2;27-43;98-104 <HAL>
A;Cross-references: UNIPARC:UPI000011EC95; UNIPARC:UPI0000171E1F; UNIPARC:UPI0000171E20
A;Cross-references: UNIPARC:UPI000011EC95; UNIPARC:UPI0000171E1F; UNIPARC:UPI0000171E20
A;Note: the sequence from the summary and from Fig. 5 is inconsistent with that from pag
A;Note: the sequence from the summary and from Fig. 5 is inconsistent with that from pag
B;Guerreiro, P.; Barreiros, T.; Soares, H.; Cyrne, L.; Maia e Silva, A.; Rodrigues-Pousa
submitted to the EMBL Data Library, April 1995
A;Description: Sequencing of a 17.6 kb segment on the right arm of yeast chromosome VII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Guerreiro, P.; Barreiros, T.; Cyrne, L.; Soares, H.; Maia
submitted to the Protein Sequence Database, May 1996
A;Reference number: S64517
                                                                                                                                                                  thioredoxin - Mycoplasma capricolum (fragment)
N;Alternate names: protein MC064
C;Species: Mycoplasma capricolum
C;Date: 09-Oct-1997 #sequence_revision 24-Oct-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yeast 12, 273-280, 1996
A;Title: Sequencing of a 17.6 kb segment on the right arm of yeast chromosome VII reveal terial electron-transferring flavoproteins (beta-chain) and of the Escherichia coli pho. A;Reference number: S63848; MUID:97060019; PMID:8904340
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A;Residues: 1-104 <SON>
A;Cross-references: UNIPARC:UPI000004F91C; EMBL:U40843; NID:g1165213; PIDN:AAA85584.1;
C;Accession: S77780; S46921
R;Bork, P.; Ouzounis, C.; Casari, G.; Schneider, R.; Sander, C.
Mol. Microbiol. 16, 955-967, 1995
A;Title: Exploring the Mycoplasma capricolum genome: a minimal
A;Reference number: S77739; MUID:96059641; PMID:7476192
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A;Experimental source: strain S288C
R;Guerreiro, P.; Barreiros, T.; Soares, H.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Gene: SGD:TRX2; TR-I; MIPS:YGR209c
A;Cross-references: SGD:S0003441; MIPS:YGR209c
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Pred. No.
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                                                                                                                   R.; Sander, C.; Dolan, M.;
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A; Molecule type: DNA
A; Residues: 1-104 <KUR>
A; Cross-references: UNIPROT: Q97P68; UNIPARC: UPI00000519A5;
A; Experimental source: strain TIGR4
C; Genetics:
                                                                                      C;Species: Streptococcus pneumoniae
C;Abte: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 09-Jul-2004
C;Abte: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 09-Jul-2004
C;Abccession: H98071
R;Hoskins, J.A.; Alborn Jr., W.; Arnold, J.; Blaszczak, L.; Burgett, S.; Dei
e, R.; LeBlanc, D.J.; Lee, L.N.; Lefkowitz, E.J.; Lu, J.; Matsushima, P.; Mc
y, P.; Sun, P.M.; Winkler, M. B.
J. Bacteriol. 183, 5709-5717, 2001
J. Bacteriol. 183, 5709-5717, 2001
A;Authors: Yang, Y.; Young-Bellido, M.; Zhao, G.; Zook, C.; Baltz, R.H.; Jas
A;Title: Genome of the Bacterium Streptococcus pneumoniae Strain R6.
A;Reference number: A97872; MUID:21429245; PMID:11544234
A;Recession: H98071
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A; Molecule type: DNA
A; Residues: 1-104 < KUR>
A; Cross-references: UNII
C; Genetics:
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H98071
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C;Superfamily:
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A;Accession: H95206
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C;Date: 03-Aug-2001 #sequence_revision 03-Aug-2001 #text_change 09-Jul-2004
C;Accession: H95206
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A;Molecule type: DNA
A;Residues: 1-104 <BOR>
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                                                                           A; Status: preliminary
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Best Local S
Matches 4
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Best Local S
Matches 4
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                  UNIPROT: Q8DNP9; UNIPARC: UPI00000E36AA;
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Pred. No. 1.
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                  GB:AE007317; PIDN:AAL00405.1
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pneumoniae.
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C46264
                                                                                                                                                                                                                                                                                                 thioredoxin 3 - slime mold (Dictyostelium discoideum)
C;Species: Dictyostelium discoideum
C;Date: 28-Oct-1994 #sequence_revision 28-Oct-1994 #t
C;Accession: C46264
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A;Contents: annotation; purification and characterization C;Superfamily: thioredoxin; thioredoxin homology
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A; Accession: A59394
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R;Reynolds, C.M.; Poole, L.B.; Hammel, K.E.; Wada, K.; Buchanan, submitted to the Protein Sequence Database, September 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: protein A; Residues: 1-104 < RE1>
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C;Date: 02-Apr-2002 #sequence_revision 02-Apr-2002
C;Accession: B59394; A59394
R;Reynolds, C.M.; Meyer, J.; Poole, L.B.
Biochemistry 41, 1990-2001, 2002
A;Title: NADH-dependent bacterial thioredoxin reduc
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C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: oxidoreductase
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Query Match
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                                                                           ;Superfamily: thioredoxin; thioredo; 8-91/Domain: thioredoxin homology
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    Clostridium pasteurianum

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Pred. No. 1.3e+02;
); Mismatches 0;
Score 31; DB 2;
Pred. No. 1.3e+02;
; Mismatches 0;
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R.Wetterauer, B.; Jacquot, J.P.; Veron, M.
J. Biol. Chem. 267, 9895-9904, 1992
A;Title: Thioredoxins from Dictyostelium discoideum are a developmentally A;Reference number: A46264; MUID:92250653; PMID:1577820
A;Accession: C46264
                                                                                                             A;Status: preliminary; not compared with conceptual translation A;Molecule type: mRNA A;Residues: 1-104 <WET> A;Residues: 1-104 <WET> A;Cross-references: UNIPROT:P29447; UNIPARC:UPI0000136D7B; GB:M91383; NID:g167932; C;Superfamily: thioredoxin; thioredoxin homology
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Nucleic Acids Res. 28, 4317-4331, 2000
A;Title: Complete genome sequence of the alkaliphilic bacterium A;Reference number: A83650; MUID:20512582; PMID:11058132
A;Accession: B84037
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-104 <870>
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B84037
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C;Accession: E89885
R;Kuroda, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.; Cui, L.; ma, A.; Mizutani-Ui, Y.; Kobayashi, N.; Sawano, T.; Inoue, R.; Kaito, C.; Sekimizu, C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K.
Lancet 357, 1225-1240, 2001
Lancet 357, 1225-1240, 2001
thioredoxin [imported] - Lactococcus lactis subsp. lactis (strain IL1403) C;Species: Lactococcus lactis subsp. lactis C;Pate: 23-Mar-2001 #sequence_revision 23-Mar-2001 #text_change 05-Oct-20 C;Pate: 23-Mar-2001 #sequence_revision 23-Mar-2001 #text_change 05-Oct-20 C;Accession: D86830 
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C;Species: Bacillus halodurans
C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 05-Oct-2004
C;Accession: B84037
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A; Residues: 1-104 < KUR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Species: Staphylococcus aureus
C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 05-Oct-2004
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C;Superfamily: thioredoxin; thioredoxin homology
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tive 0; Mismatches 0;
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Pred. No. 1.3e+02;
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                                                                       #text_change 05-Oct-2004
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RESULT 37
C81432
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R;Parkhill, J.; Wren, B.W.; Mungall, K.; Ketley, J.M.; Churcher, C.; Basham, D.; Chillin C.W.; Quail, M.; Rajandream, M.A.; Rutherford, K.M.; VanVliet, A.; Whitehead, S.; Barrel Nature 403, 665-668, 2000
A;Title: The genome sequence of the food-borne pathogen Campylobacter jejuni reveals hyla;Reference number: A81250; MUID:20150912; PMID:10688204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Title: The complete genome sequence of the lactic acid bacterium Lactococcus lactis A;Reference number: A86625; MUID:21235186; PMID:11337471 A;Accession: D86830 A;Status: preliminary
                                                             A;Reference number: A30006; MUID:88257080; PMID:2838473
A;Accession: A30006
A;Molecule type: mRNA
A;Residues: 1-105 <JON>
A;Cross-references: UNIPROT:P08629; UNIPARC:UPI00001713BC;
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
E;9-92/Domain: thioredoxin homology <THR>
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A;Cross-references: UNIPROT:Q9CF37; UNIPARC:UPI00000C6AAD;
                                                                                                                                                                                                                        Title: Isolation of a chicken thioredoxin cDNA clone: A;Title: Isolation of a chicken thioredoxin cDNA clone: A;Title: A;Title: Isolation of a chicken thioredoxin cDNA clone:
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C;Superfamily: thioredoxin; thioredoxin homology
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A; Residues: 1-104 < P
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C;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 05-Oct-2004
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C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                          C;Species: Gallus gallus (chicken)
C;Date: 31-Mar-1989 #sequence_revision
                                                                                                                                                                                                                                                                                                                                                                                            thioredoxin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Experimental source: strain IL1403
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                                             F;32-35/Disulfide bonds:
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100.0%; E
tive 0;
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100.0%;

Score Pred.

NO ;

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DB 1;

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A; Molecule type: mRNA
A; Residues: 1-105 < TAG2>
A; Cross-references: UNIPARC: UPI000011065C; EMBL: X77584; NID: 9453963; PIDN: CAA54687.1; PIR; Wollman, E.E.; d'Auriol, L.; Rimeky, L.; Shaw, A.; Jacquot, J.P.; Wingfield, P.; Grabel, Biol. Chem. 263, 15506-15512, 1988
A; Title: Cloning and expression of a cDNA for human thioredoxin.
A; Reference number: A31993; MUID: 89008454; PMID: 3170595
A; Accession: A31993
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J. Biol. Chem. 267, 9835-9904, 1992
A;Ticle: Thioredoxins from Dictyostellum discoideum are
A;Reference number: A46264; MUID:92250653; PMID:1577820
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                                                                                                                                                                                                                      A;Reference number: S44375; MUID:94244626; PMID:8187776
A;Contents: erratum
                                                                                                                                                                                                                                                                    A;Cross-references: UNIPARC:UPI000011065C; GB:X77584; NID:g453963; PIDN:CAA54687.1; PIDA;Note: this sequence has been revised in reference S44375
R;Pagaya, Y.; Maeda, Y.; Mitsui, A.; Kando, N.; Matsui, H.; Hamuro, J.; Brown, N.; Arai
                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Title: ATL-derived factor (ADF), an IL-2 receptor/Tac
A;Reference number: S04106; MUID:89251607; PMID:2785919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:P10599; UNIPARC:UPI000011065C; EMBL:X54539; NID:g37455; PIDNR;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kondo, N.; Matsui, H.; Hamuro, J.; Brown, N.; Arai, EMBO J. 8, 757-764, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gene 102, 221-228, 1991
A; Title: Isolation and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: mRNA
A; Residues: 1-105 < WE'
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A;Status: prelimina
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A;Residues: 1-105 <TAG1>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ,....e: 1801ation and characterization of human thioredoxin-encoding genes;Reference number: JH0568; MUID:91340156; PMID:1874447;RACCESSION: JH0568
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;Date: 30-Jun-1992 #sequence revision 26-May-1994 #text change 05-Oct-2004;Accession: JH0568; S04106; S44375; A31993; PT0079; A60749; A38922; S53453;;Tonissen, K.F.; Wells, J.R.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Accession: S04106
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A;Contents: annotation; X-ray crystallography, 2.1 angstroms, oxidized for R;Forman-Kay, J.D.; Clore, G.M.; Gronenborn, A.M. submitted to the Brookhaven Protein Data Bank, December 1990 A;Reference number: A50924; PDB:4TRX A;Referents: annotation; conformation by (1)H-NMR, residues 1-73,'T',75-105 R;Forman-Kay, J.D.; Clore, G.M.; Wingfield, P.T.; Gronenborn, A.M. Biochemistry 30, 2685-2698, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Note: described to be a surface-associated thioredoxin R;Wakasugi, H.; Rimsky, L.; Mahe, Y.; Kamel, A.M.; Fradelizi, I Proc. Natl. Acad. Sci. U.S.A. 84, 804-808, 1987
A;Title: Epstein-Barr virus-containing B-cell line produces an A;Reference number: A60870; MUID:87118252; PMID:3027706
A;Contents: annotation R;Weichsel, A.; Gasdaska, J.R.; Powis, G.; Montfort, W.R. submitted to the Brockhaven Protein Data Bank, February 1996
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R;Silberstein, D.S.; Ali, M.H.; Baker, S.L.; David, J.R.
J. Immunol. 143, 979-983, 1989
A;Title: Human eosinophil cytotoxicity-enhancing factor.
A;Reference number: A60749; MUID:89309777; PMID:2745979
                                                                                                                                                                                                                         A;Map position: 9q31-9q31
A;Introns: 8/3; 43/3; 85/3
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Contents: annotation; conformation by (1)H- and C; Comment: This small ubiquitous protein functions C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: High-resolution three-dimensional structure of red A;Reference number: A38953; MUID:91159399; PMID:2001356 A;Contents: annotation; conformation by (1)H- and (15)N-NMR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Contents: annotation; X-ray crystallography, 1.7 angstroms, R;Weichsel, A.; Gasdaska, J.R.; Powis, G.; Montfort, W.R. submitted to the Brookhaven Protein Data Bank, February 1996
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Biochem. J. 304, 861-867, 1994
A;Title: Characterization of a thioredoxin-related surface protein.
A;Reference number: SS3453; MUID:95118305; PMID:7818492
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J. Immunol. 136, 3304-3310, 1986
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A;Molecule type: protein
A;Residues: 2-12, 'K',14-15,'XX',18-19,'X',21-22 <SIL>
A;Cross-references: UNIPARC:UPI0000171E1A
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Biochem. Biophys. Res. Commun. 175, 123-128, 1991
A;Title: Identification of a thioredoxin-related protein associated with plasma membran A;Reference number: PT0079; MUID:91151337; PMID:1998498
                                                                                                                                                                                                                                                                                                                                                                           A;Gene: GDB:TXN
A;Cross-references: GDB:120475; OMIM:187700
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A;Residues: 1-21;38-57 <DEA>
A;Cross-references: UNIPARC:UPI00000353C5; UNIPARC:UPI0000171E1C
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A;Cross-references: UNIPARC:UPI000003066B; GB:J04026; NID:g339648; PIDN:AAA74596.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Molecule type: protein
A;Residues: 2-16 <WAK>
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A;Residues: 2-13,'X',15 <MAR>
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                                                                                                                  F;32-35/Disulfide
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                                                                                                                                                                                     :Reywords: redox-active disulfide
:2-105/Product: thioredoxin #status experimental <MAT>
Query Match
Best Local Similarity
                                                                                                                                                      9-92/Domain: thioredoxin homology
                                                                                                                  bonds:
                                                                                                              redox-active #status experimental
100.0%;
Score 31;
Pred. No.
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1.3e+02;

human thiore

path

residues

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C;Species: Macaca
C;Date: 30-Jun-19
C;Accession: JS06
R;An, G.; Wu, R.
Biochem. Biophys.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Species: Mus musculus (house mouse)
C;Date: 21-Nov-1993 #sequence revision 17-Oct-1997 #text_change 05-Oct-2004
C;Accession: JC4068; S44376; S304107
R;Matsul, M.; Taniguchi, Y.; Hirota, K.; Taketo, M.; Yodoi, J.
Gene 152, 165-171, 1995
A;Title: Structure of the mouse thioredoxin-encoding gene and its processed A;Reference number: JC4068; MUID:95137382; PMID:7835695
A;Accession: JC4068
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JS0667
A;Map position: 4:24.6
A;Introns: 29/2; 44/1;
C;Superfamily: thioredo
                                                                                                                                                                                                                            A;Cross-references: UNIPARC:UPI0000163891; EMBL:X77585; NID:g453971; PIDN:CAA54688.1; R;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kondo, N.; Matsui, H.; Hamuro, J.; Brown, N.; Ara EMBO J. 8, 757-764, 1989
A;Title: ATL-derived factor (ADF), an IL-2 receptor/Tac inducer homologous to thioredo A;Reference number: S04106; MUID:89251607; PMID:2785919
A;Accession: S04107
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S04107
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A;Title: Thioredoxin gene expression is transcriptionally up-regulated by retinol
A;Reference number: JS0667; MUID:92181438; PMID:1543487
A;Accession: JS0667
                                                                                                                                 A;Molecule type: mRNA
A;Residues: 1-93,'N',94-96,'ALT',100-104,'S' <TAG2>
A;Residues: UNIPARC:UPI0000171E1E; GB:X77585
A;Cromsment: This small ubiquitous protein functions in
                                                                                                                                                                                                                                                                                                                                                                                                               A;Reference number: S44375; MUID:94244626; PMID:8187776
A;Contents: erratum
A;Accession: S44376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:P10639; UNIPARC:UPI0000163891; DDBJ:D21855; NID:g517128 R;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kando, N.; Matsui, H.; Hamuro, J.; Brown, N.; Arai EMBO J. 13, 2244, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-105 < MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N;Alternate names: ATL-derived factor (ADF)
C;Species: Mus musculus (house mouse)
C;Date: 21-Nov-1993 #sequence revision 17-C
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                                                                  A; Cross-references: MGI:36258
                                                                                             A;Gene: MGI:Txn
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A;Residues: 1-105 <TAG1>
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;Date: 30-Jun-1992 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
;Accession: JS0667
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Best Local
                                                                                                                  Genetics:
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  thioredoxin; thioredoxin homology
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C;Accession: A00281
R;Meng, M.; Hogenkamp, H.P.C.
J. Biol. Chem. 256, 9174-9182, 1981
A;Title: Purification, characterization, an
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A;Title: Characterization of a thioredoxin-related surface protein. A;Reference number: S53453; MUID:95118305; PMID:7818492
                                                                       A;Molecule type: protein
A;Residues: 1-105 <MEN>
A;Residues: 1-105 <MEN>
A;Residues: 1-105 <MEN>
A;Cross-references: UNIPACT:P00275; UNIPARC:UPI0000136D69
A;Cross-references: UNIPACT:P00275; UNIPARC:UPI0000136D69
C;Comment: Thioredoxins are ubiquitous small hydrogen carrier proteins that participate C;Cuperfamily: thioredoxin; thioredoxin homology
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A;Note: 12-Lys, 15-Gly, 16-Leu, 17-Gln, and 18-Leu were also
A;Note: described to be a surface-associated thioredoxin
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 31-Mar-1990 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
C;Accession: S04352; S66372
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                         C; Keywords: redox-active disulfide F;8-91/Domain: thioredoxin homology <THR>
                                                                                                                                                                                                                                                                                                                                                                                                         thioredoxin - coryneform bacterium ATCC11425
C;Species: coryneform bacterium ATCC11425
C;Date: 02-Apr-1982 #sequence_revision 02-Apr-1982 #text_change 05-Oct-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: protein
A; Residues: 2-21 <DEA>
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                                                                                                                                                                                                                                              A; Reference number: A00281; A; Accession: A00281
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A; Residues: 1-105 < TON>
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;30-33/Disulfide bonds: redox-active
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#status experimental
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5; PMID:7021558
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R;Ogata, H.; Audic, S.; Renesto-Audiffren, P.; Fournier, P.E.; Barbe, V.; Samson, Science 293, 2093-2098, 2001
A;Title: Mechanisms of Evolution in Rickettsia conorii and Rickettsia prowazekii.
A;Reference number: A97700; MUID:21442074; PMID:11557893
A;Accession: B97700
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT:P52232; UNIPARC:UPI0000136D6E; EMBL:D64000; GB:AB001339; NIIA;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1996 C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Synechocystis sp.
A;Variety: PCC 6803
C;Date: 25-Apr-1997 #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004
C;Accession: $76386
C;Accession: $76386
S; Kotani, H.; Tanaka, A.; Asamizu, E.; Nakamura, Y.; l
o, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamado
DNA Res. 3, 109-136, 1996
RESULT
AG2042
                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Rickettsia conorii
C;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 05-Oct-2004
C;Accession: B97700
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A; Residues: 1-105 < KUR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Title: Sequence analysis of the genome of the unicellular cyanobacterium Synechocysti
                                                                                                                                     Query Match
Best Local S
Matches 4
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100.0%; Pred. No. 1.3e+02;
. Mismatches 0;
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100.0%; Pred. No. 1.3e+02;
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Pred. No.
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Yamada, M.; Yasud
                                                                                                                                                                                                                                                              PIDN:AAL02540.1
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R;Pille, S.; Chuat, J.C.; Breton, A.M.; Cle
J. Bacteriol. 172, 1556-1561, 1990
A;Title: Cloning, nucleotide sequence, and
A;Reference number: A35135; MUID:90170874;
A;Accession: A35135
                                                                                                                                                                            C;Species: Rhodobacter sphaeroides
C;Date: 27-Jul-1990 #sequence_revision
C;Accession: A35135; A30508
                                                                                                                                                                                                                                                                      RESULT
A35135
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J. Bacteriol. 183, 4823-4838, 2001
A,Title: Genome Sequence and Comparative Analysis of the Solvent-Producing Bacterium
A:Reference number: A96900; MUID:21359325; PMID:21359325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tnioredoxin [imported] - Clostridium acetobutylicum
C;Species: Clostridium acetobutylicum
C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 05-Oct-2004
C;Accession: D97279
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A;Cross-references: UNIPROT:Q8YVT1; UNIPARC:UPI00000CE247; GB:BA000019; PIDN:BAB73592.1
A:Experimental source: strain PCC 7120
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A;Note: Nostoc sp. strain PCC 7120 is a synonym of
C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001
C;Accession: AG2042
A;Molecule type: DNA
A;Residues: 1-106 <PII>
A;Cross-references: UNIPROT:P08058; UNIPARC:UPI000016FF78; GB:M33806; NID:g152044;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Gene: CAC3083
C; Superfamily: thioredoxin; thioredoxin homology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Gene: all1893
C;Superfamily:
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A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium An
A;Reference number: AB1807; MUID:21595285; PMID:11759840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:Q97EM7; UNIPARC:UPI00000CA715; GB:AE001437; PIDN:AAK81023.1,
A;Experimental source: Clostridium acetobutylicum ATCC824
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A; Residues: 1-105 < KUR>
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A;Status: preliminary
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Best Local Similarity
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Best Local
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    Rhodobacter sphaeroides

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                                                                                                                                                          A.M.; Clement-Metral, J.D.; Galibert,
                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 31; DB 2;
Pred. No. 1.3e+02
); Mismatches 0
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Pred. No. 1.3e+02;
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PMID:2137818
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#text_change 05-Oct-2004
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Taylor,

D.E.; Vovis, G.F.;

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R;Clement-Metral, J.D.; Holmgren, A.; Cambillau, C.; Joernvall, H.; I Eur. J. Biochem. 172, 413-419, 1988
A;Title: Amino acid sequence determination and three-dimensional mod. A;Reference number: A30508; MUID:88166714; PMID:3280308
A;Accession: A30508
A;Molecule type: protein
A;Cross-references: UNIPARC:UPI0000171E23
A;Cross-references: UNIPARC:UPI0000171E23
A;Experimental source: strain Y
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
F;9-92/Domain: thioredoxin; thoology
F;31-34/Disulfide bonds: redox-active #status predicted
                                                                                                         C;Accession: H64622; C71890

R;Tomb, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; Fleischmann, Peterson, S.; Loftus, B.; Richardson, D.; Dodson, R.; Khalak, H.G.; Glodek, A.; Mcison, J.D.; Kelley, J.M.; Cotton, M.D.; Weidman, J.M.; Fujii, C.; Bowman, C.; Watthey Nature 388, 539-547, 1997

A;Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser A;Title: The complete genome sequence of the gastric pathogen Helicobacter pylori. A;Reference number: A64520; MUID:97394467; PMID:9252185

A;Accession: H64622
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 A;Experimental source: strain 26695 R;Alm, R.A.; Ling, L.S.L.; Moir, D.
                                                                                                                                                                                                                                                                                                      C; Species: Helicobacter pylori
A; Variety: strains J99, 26695
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A; Accession: $33357
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R;Labudova, O.; Nemethova,
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                                                        A; Molecule type: DNA
A; Residues: 1-106 < TOM>
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                                     A;Cross-references: UNIPROT:P56430; UNIPARC:UPI0000136E82;
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Matches 4
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;9-92/Domain: thioredoxin homology <THR>

;31-34/Disulfide bonds: redox-active #status
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;Residues: 1-106 <LAB>
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;pecies: Streptomyces aureofaciens
;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change
;Accession: S33357
                                                                                            Status: nucleic acid sequence
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Library, April 1993
of a protein-coding part
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 A; Experimental source: C; Genetics:
                   A;Cross-references: UNIPROT:Q8UJA6; UNIPARC:UPI0000164451; GB:AE008688; PIDN:AAL41053.1 A;Experimental source: strain C58 (Dupont)
                                                      A; Molecule type: DNA
A: Residues: 1-106 < KUR>
                                                                                                                                                   ster, E.W.
A; Title: The Genome of
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A; Residues: 1-106 < COH>
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                                                                                            A;Status: preliminary
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A;Accession: AG2579
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C;Accession: AG2579
R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, Y.; Woo, erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, M.; McClel; Karp, P.; Romero, P.; Zhang, S.
Science 294, 2317-2323, 2001
A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.; Gordon-Kamm,
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                                                                                                                                                                                                                                        thioredoxin C-1 trxA [imported] - Agrobacterium tumefaciens (strain C58, Dupont) C;Species: Agrobacterium tumefaciens C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 05-Oct-2004
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A;Title: The thioredoxin system of Penicillium chrysogenum and its p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Superfamily: thioredoxin; thioredoxin homology F;10-92/Domain: thioredoxin homology <THR>
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of the Natural Genetic AB2577; MUID:21608550;
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Pred. No.
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Engineer Agrobacterium tumefaciens PMID:11743193
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PMID:9923682
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A;Cross-references: UNIPARC:UPI00001655B5; GB:M14736; NID:g142117; PIDN:AAA22049.1;
R;Gleason, F.K.; Whittaker, M.M.; Holmgren, A.; Joernvall, H.

J. Biol. Chem. 260, 9567-9573, 1985
A;Title: The primary structure of thioredoxin from the filamentous cyanobacterium An A;Reference number: A23910; MUID:85261357; PMID:3926769
A;Reference number: A23910; MUID:85261357; PMID:3926769
A;Residues: 2-107 <GLE>
A;Residues: 2-107 <GLE>
A;Residues: 2-107 <GLE>
A;Residues: 2-107 <GLE>
A;Stross-references: UNIPARC:UPI0000136D66
A;Stross-references: UNIPARC:UPI0000136D66
A;Stross-references: UNIPARC:UPI0000136D66
A;Bxperimental source: PCC 7119, ATCC 29151
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: heat-stable protein; redox-active disulfide
F;22-107/Product: thioredoxin homology <THR>
F;10-93/Domain: thioredoxin homology <THR>
F;32-35/Disulfide bonds: redox-active #status predicted
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C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
F;10-93/Domain: thioredoxin homology <THR>
F;32-35/Disulfide bonds: redox-active #status pr
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J. Biol. Chem. 264, 4008-4014, 1989
A;Title: Thioredoxin is essential for photosynthetic growth.
A;Reference number: A32956; MUID:89139466; PMID:2492995
A;Accession: A32956
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A32956
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C;Species: Synechococcus sp.
C;Date: 17-Jul-1992 #sequence revision
C;Accession: A32956; A30842
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A;Map position: circular chromosome
C;Superfamily: thioredoxin; thioredo
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A; Residues: 1-107 < RES>
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A; Residues: 1-107 < MUL>
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Dec.1988 #sequence_revision 11-Apr-1997 #text_change 05-Oct-2004;Date: 31-Dec.1988 #sequence_revision 11-Apr-1997 #text_change 05-Oct-2004
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Pred. No. 1.3e+02;
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A,Cross-references: UNIPROT:P37395; UNIPARC:UPI0000136E7B; EMBL R;Ohta, N.; Kawano, S.; Kuroiwa, T. Curr. Genet. 26, 136-138, 1994.
A;Title: Physical map of the plastid genome of the unicellular A;Reference number: $45488; MUID:95094309; PMID:8001167
A;Accession: $45488
                                                                          A;Gene: trx
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
C;Keywords: redox-active disulfide
F;10-93/Domain: thioredoxin homology <THR>
F;32-35/Disulfide bonds: redox-active #status pr
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C,Superfamily: thioredoxin; thioredoxin homology
C;Keywords: heat-stable protein; redox-active disulfide
F;10-93/Domain: thioredoxin homology <THR>
                                                                                                                                                                                              A;Molecule type: DNA
A;Residues: 'M',6,'IDEL',11-12,'DT',15-16,'LQ',19,'HQ',22-38,'A',40,'IL',43-44,'I',46-4
A;Cross-references: UNIPARC:UPI00000A275B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R;Langsdorf, A.; Emich, A.; Zetsche, K. submitted to the EMBL Data Library, February A;Description: A thioredoxin gene is located
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: protein A; Residues: 1-107 < JOH>
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A;Title: The primary structure of thioredoxin from Chromatium vinosum A;Reference number: A26622; MUID:87185419; PMID:3567166
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C;Date: 31-Mar-1988 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
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A; Residues: 1-107 < LAN>
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C;Species: Cyanidium caldarium
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A;Cross-references: UNIPROT:P09857; UNIPARC:UPI0000136E76
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              100.0%; Score 31; DB 1; 1 100.0%; Pred. No. 1.3e+02;
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thioredoxin A - red alga (Porphyra purpurea) chloroplast
C;Species: chloroplast Porphyra purpurea
C;Date: 19-Mar-1997 #sequence_revision 09-May-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Salz, H.K.; Plickinger, T.W.; Mittendorf, E.; Pellicena-Palle, A.; Petschek, J.P.; Brogenetics 136, 1075-1086, 1994
A;Title: The Drosophila maternal effect locus deadhead encodes a thioredoxin homolog req A;Reference number: S47867; MUID:94274010; PMID:7516301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 thioredoxin-like protein - fruit fly (Drosophila melanogaster) C;Species: Drosophila melanogaster C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change C;Accession: S47867
                       RESULT 60
S46521
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S73146
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A; Residues: 1-107 < SAL>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Plant Mol. Biol. Rep. 13, 333-335, 1995
A;Title: Complete nucleotide sequence of the Porphyra purpurea chloroplast genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C; Accession: S73146
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  thioredoxin -
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Best Local S
Matches 4
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;8-92/Domain: thioredoxin homology <THR>
;31-34/Disulfide bonds: redox-active #status predicted
                                                                                                                                                                                                                                                                                Superfamily: thioredoxin; thioredoxin homology; Keywords: chloroplast; redox-active disulfide; 9-92/Domain: thioredoxin homology <THR>
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Porphyra yezoensis chloroplast
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Pred. No.
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Pred. No.
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  A;Molecule type: DNA
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C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #LEAK. C;Accession: T33843
R;Woessner, J.; Graves, T.; Keppler, D.
submitted to the EMBL Data Library, November 1998
A;Description: The sequence of C. elegans cosmid Y44E3A.
A;Reference number: Z21422
                                                                                       C;Species: Synechocystis sp.
A;Variety: PCC 6803
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995
C;Accession: S46958; S76679
R;Navarro, F; Florencio, FJ.
                                                                                                                                                                                          thioredoxin A - Symechocystis sp. N;Alternate names: protein slr0623 C;Species: Synechocystis sp.
                                                                                                                                                                                                                                                                   RESULT 62
S46958
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A;Introns: 31/2
C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Molecule type: DNA
A;Residues: 1-107 <WOB>
A;Cross-references: UNIPROT:Q9TXY8; UNIPARC:UPI0000077021; EMBL:AF106589; PIDN:AAC78230
A;Experimental source: strain Bristol N2; clone Y44E3A
C;Genetics:
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C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: chloroplast; redox-active disulfide
F;9-92/Domain: thioredoxin homology <THR>
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A;Residues: 1-107 <REY'S
A;Cross-references: UNIPROT:P50254; UNIPARC:UPI0000136E94; EMBL:X76612; NID:g509276; PIC
C;Genetics:
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Plant Mol. Biol. 25, 13-21, 1994
A;Title: Chloroplast encoded thioredoxin genes in the red algae Porphyra yezoensis and A;Reference number: S46521; MUID:94272009; PMID:8003693
A;Accession: S46521
A;Status: preliminary
A;Status: preliminary
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C;Date: 19-Mar-1997 #sequence_revision 06-Jun-1997 #text_change 09-Jul-2004
C;Accession: S46521
submitted to the EMBL Data Library, July 1994 A;Description: The gene encoding for thioredo: A;Reference number: S46957 A;Accession: S46958
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Best Local (
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Best Local Similarity
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Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change
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100.0%; Pred. No. 1.3e+02;
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Pred. No. 1.3e+02;
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thioredoxin - Streptomyces clavuligerus
C;Species: Streptomyces clavuligerus
C;Date: 08-Sep-1995 #sequence_revision 08-Sep-1995 #text_change 05-Oct-2004
C;Accession: B53307
R;Cohen, G; Yanko, M; Mislovati, M; Argaman, A; Schreiber, R; Av-Gay, Y;; I
J. Bacteriol. 175, 5159-5167, 1993
A;Title: Thioredoxin-thioredoxin reductase system of Streptomyces clavuligerus: A;Reference number: A53307; MUID:93352422; PMID:8349555
A;Accession: B53307
A;Status: preliminary
A;Molecule type: DNA
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A; Map position:
C; Superfamily:
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C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
F;10-93/Domain: thioredoxin homology <THR>
F;32-35/Disulfide bonds: redox-active #status predicted
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B53307
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A;Residues: 1-107 <NAV>
A;Cross-references: UNIFROT:P52231; UNIPARC:UPI0000164C7C; EMBL:X80486; NID:g515939; PII R;Kaneko, T.; Sato, S.; Kotani, H.; Tanaka, A.; Asamizu, E.; Nakamura, Y.; Miyajima, N.; R;Kaneko, T.; Sato, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamada, M.; Yasuda DNA Res. 3, 109-136, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Date: 24-Mar-1999 #sequence_revision 24-Mar-1999 #text_change 09-Jul-2004 C;Accession: B81458; T02814 R;Myler, P.J.; Budleman, L.; deVos, T.; Hixson, G.; Kiser, P.; Lemley, C.; Magness, C.; Proc. Natl. Acad. Sci. U.S.A. 96, 2902-2906, 1999 A;Title: Leishmania major Friedlin chromosome 1 has an unusual distribution of protein-A;Reference number: A81455; MUID:99178987; PMID:10077609
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A; Residues: 1-107 < K
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Matches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Status: nucleic acid sequence not shown; translation not
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A;Residues: 1-107 <COH>
A;Cross-references: UNIC;Genetics:
                                                                                                                                                                                                                                    R;DelVecchio, V.G.; Kapatral, V.; Redkar, R.J.; Patra, G.; Mujer, C.; Los, T.; Ivanova, ; Mazur, M.; Goltsman, E.; Selkov, E.; Elzer, P.H.; Hagius, S.; O'Callaghan, D.; Letess Proc. Natl. Acad. Sci. U.S.A. 99, 443-448, 2002
A;Title: The genome sequence of the facultative intracellular pathogen Brucella melitens A;Reference number: AD3252; PMID:11756688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Species: Nostoc sp. PCC 7120
A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. strain PCC 7120
C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change 05-Oct-2004
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C;Superfamily: thioredoxin;
F;11-93/Domain: thioredoxin
                                                                 A; Map position:
C; Superfamily:
                                                                                                                                   A; Cross-references: UNIPROT: Q8YE56; A; Experimental source: strain 16M
                                                                                                                                                                                                                                                                                                                                      C;Species: Brucella melitensis
C;Date: 01-Feb-2002 #sequence_revision 01-Feb-2002 #text_change 05-Oct-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Gene: trxA
C; Superfamily:
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A; Residues: 1-107 < KUR>
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A; Title: Complete Genomic
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                                                                 ;Superfamily: thioredoxin; thioredoxin homology
               Query Match
Best Local S
                                                                                                   Gene:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     , Y.; Wolk, C.P.; Kuritz, T.;
S.; Sugimoto, M.; Takazawa, N
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7; MUID:21595285; PMID:11759840
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Pred. No.
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M.; Yamada,
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Similarity 4; Conserv

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R;Fleischmann, R.D.; Adams, M.D.; White, O.; Clayton, R.A.; Kirkness, E.P.; Kerlavage, Gocayne, J.D.; Scott, J.; Shirley, R.; Liu, L.I.; Glodek, A.; Kelley, J.M.; Weidman, D.M.; Brandon, R.C.; Fine, L.D.; Fritchman, J.L.; Fuhrmann, J.L.; Geoghagen, N.S.M.; Science 269, 496-512, 1995
A;Authors: Gnehm, C.L.; McDonald, L.A.; Small, K.V.; Fraser, C.M.; Smith, H.O.; Venter, A;Title: Whole-genome random sequencing and assembly of Haemophilus influenzae Rd. A;Reference number: A64000; MUID:95350630; PMID:7542800
A;Accession: E64047
                                                                                                                                                                                                                                                                                                                                                                   A; Status. F-2--A; Molecule type: DNA A; Molecule type: DNA A; Residues: 1-108 <STO>A; Residues: 1-108 <STO>A; Cross-references: UNIPROT: Q9X2T1; UNIPARC: UPI0000136E95; GB: AE004936; GB: AE004091; A; Cross-references: Strain PAO1
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C;Date: 18-Aug-1995 #sequence_revision 18-Aug-1995 #text_change 05-Oct-2004
C;Accession: E64047
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A;Accession: G82991
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R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig,
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  thioredoxin -
                            RESULT 69
B55124
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Nature 406, 959-964, 2000
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C;Species: Pseudomonas aeruginosa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Status: preliminary
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A;Experimental source: strain Tassajara
A;Note: unidentified residues are Ile or Leu
A;Note: the species is identified as Chlorobium thiosulfatophilum
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: photosynthesis; redox-active disulfide
F;10-93/Domain: thioredoxin homology cTHR>
F;32-35/Disulfide bonds: redox-active #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: UNIPARC:UPI0000174DEC
A;Note: unidentified residues are IIe or Leu
A;Note: the species is identified as Chlorobium thiosulfatophilum
R;Mathews, W.R.; Johnson, R.S.; Cornwell, K.L.; Johnson, T.C.; Bud
J. Biol. Chem. 262, 7537-7545, 1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Gene: trxA
C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                                                                                                      R;Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; Prentice, deno-Tarraga, A.M.; Chillingworth, T.; Cronin, A.; Davies, R.M.; Davis, P.; Dougan, il, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; Whitehead, S.; Barre Nature 413, 523-527, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   thioredoxin 1 [imported] - Yersinia pestis (strain CO92)
C;Species: Yersinia pestis
C;Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 05-Oct-2004
C;Accession: AD0471
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A;Residues: 1-15,'XX',18-41,'X',43-44,'X',46-53,'XX',56-65,'X',67-74,'X',76-78,'XX',81-5
A;Cross-references: UNIPARC:UPI0000174DBD
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C;Genetics:
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A; Residues: 1-108 < KUR>
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A;Accession: B55124
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Q733M6—BACC1 Q81AFB BACCR Q81Y83—BACHA Q81Y83—BACHK Q6371L5—BACCZ Q9YEH7—AERPE Q6YQ14—HYDMA Q66NW9 RHILE	TRXF MESCR Q9NB50 RIFPA Q9SNB5 MACFA Q9SNB5 BACCF Q4MNH0 BACCE Q8DQ10 STRR6	VEGFA BOVIN VEGFA HORSE VEGFA PIG 077643 SHEEP	Q655X0 ORYSA Q8H2V6 ORYSA Q66PE7 9BRAD TRXF SPICE	Q8R2V7_MOUSE Q8R2V7_HOUSE Q96MK4_HUMAN O95LO4_FELCA	Q6ZVAZ_HUMAN Q8S091_ORYSA Q8DTZ1_STRMU	Q66PC7_RHILE O50189_MYCHY Q62GX6_BURMA	Q8LG26_ARATH Q56YG4_ARATH Q66PC4_RHILE	Q97R36_STRPN Q9RXW6_DEIRA TRXM2_ARATH	TF2LY HUMAN TRXF2 ARATH O9MZJ6 MACMU	Q81S90_BACAN Q6HKG7_BACHK Q63D26_BACCZ	Q4MUIO_BACCE Q73AC8_BACC1 Q81F76_BACCR	Q66PE6_9BRAD Q8HY70_MUSVI Q4NOC9_9DELT	Q8LFA3 ARATH Q9M7X9 ARATH O66PE1 RHIME	TRXX_ARATH Q7Z4P9_HUMAN	Q4SS63 TETNG TRXF_BRANA	Q6JE37_NICBE	Q6QAT5_DROSI	TRXM SPIOL	Q66PDO_RHILE	Q95AH9 PEA	TRXM1 ARATH Q9A3D7 CAUCR	OB3WD6 MICGR
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InterPro; IPR007958; toxin 5.
Pfam; PF05294; Toxin 5; 1.
Direct protein sequencing; Neurotoxin; Toxin.
DISULFID 1 18 By similarity.
DISULFID 4 25 By similarity.
DISULFID 15 30 By similarity.
DISULFID 19 32 By similarity.
SEQUENCE 35 AA; 3673 MW; 213E69262289EB5A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Androctonus mauretanicus mauretanicus (Scorpion).
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida;
Buthida; Buthoidea; Buthidae; Androctonus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-JUL-1986 (Rel.
21-JUL-1986 (Rel.
05-JUL-2004 (Rel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rosso J.-P., Rochat H.;
"Characterization of ten proteins
scorpion Androctonus mauretanicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=6860;
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                                                                                                                                                                                                                                                                                                                      PIR; A01758; NTSRPM.
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TISSUE SPECIFICITY: Expressed by the venom gland.
SIMILARITY: Belongs to the short scorpion toxin family.
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Q93NF4_ARTNI
Q92234_RAT
Q8NABO_HUMAN
Q6UWD8_HUMAN
Q7K4P2_DROME
Q8C708_MOUSE
Q8C708_MOUSE
Q6P3A6_MOUSE
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Q6SN76_9PR.IM
Q6SNH6_COLGU
Q6SNP3_PAPPA
Q8LDI1_ARATH
Q6ASV9_PROAC
Q8RG01_FUSIN
Q9CB61_MYCLE
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Pred. No.
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                                             Mismatches
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Voelter W.
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14 BUTSI STANDARD; PRT; 36

P59887;
10-OCT-2003 (Rel. 42, Created)
10-OCT-2003 (Rel. 42, Last sequence updat
05-UUL-2004 (Rel. 44, Last annotation upc
Chlorotoxin-like peptide Bs 14 (Bs14).
Buthus sindicus (Scorpion).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          from scorpion (Buthus sindicus) venom.";
Comp. Biochem. Physiol. 121A:323-332(1998).
-!- FUNCTION: Probable chloride channel lig.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
Buthida; Buthoidea;
                                       Name=LH-beta;
Tapirus indicus (Asiatic tapir) (Malayan tapir).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Laurasiatheria; Perissodactyla;
                                                                                                                                                                                           046621_TAPIN PRELIMINARY;
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MEDLINE=99157745;
      Tapirus.
NCBI_TaxID=9802;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conic channel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           conductance chloride channels (By similarity).
SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Expressed by the venom gland
MASS SPECTROMETRY: MW=3848.3; METHOD=MALDI; RANG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NOTE=Ref.1.
SIMILARITY: Belongs to the s
Channel inhibitor subfamily.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                European Bioinformatics Institute.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PF05294; Toxin 5; 1.

ide channel inhibitor; Direct protein
                                                                                                                                                                                                                                                                                                                                                               Similarity
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ilarity 100.
Conservative
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                                                                                                             (TIEMBLIE1. 06, Created)
(TIEMBLIE1. 06, Last sequence update)
(TIEMBLIE1. 23, Last annotation update)
hormone beta-subunit (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                              il inhibitor; Neurotoxin; 18 By simila 4 By simila 4 30 By simila 15 30 By simila 19 32 By simila
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S., Schuetz J., Kayed R., Abbasi A., Zaidi Z.H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AND MASS
                                                                                                                                                                                                                                                                                                                                                                                                                 3847
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Buthidae; Mesobuthus.
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By similarity.
By similarity.
S36117F8E50165A0
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hes 0;
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                                                 Euteleostomi;
Tapiridae;
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RESULT 4

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ID 046620 E

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DT (Fragmen

GN Name=LH/

OS Equus as

OC Eukaryot

OC Eukaryot

OC Eukaryot

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OC Mammalia

OC EUKaryot

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Best Local (
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947_DROHY
Q23947_DROHY PRELIMINARY;
Q23947,
Q23947,
Q23947,
Q23947,
Q1-NOV-1996 (TremBirel. 01, C
01-NOV-1996 (TremBirel. 26, L
01-MAR-2004 (TremBirel. 26, L
Testis-specific RNA.
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01-JUN-1998
01-JUN-2003
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NON_TER
SEQUENCE
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EMBL; AF04766; AAC04364.1; -; Genomic_DNA.
HSSP; P01233; 1HCN.
GO; GO: Onnerra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE.
Fischer S., Veits J., Meyer H.H.D.;
Fischer S., Veits J., Meyer H.H.D.;
Submitted (PEB-1998) to the EMBL/GenBank/DD
EMBL; AF047605; AAC04363.1; -; Genomic_DNA.
HSSP; P01233; 1HCN.
                                                            Name=Mst87Fa; Synonyms=DhTc3;
Drosophila hydei (Fruit fly)
Eukaryota; Metazoa; Arthropoda; Hexapoda; Inse
Neoptera; Endopterygota; Diptera; Brachycera;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GO; GO:0005179; F:hormone activity; IEA.
InterPro; IPR006208; Cys_knot.
Pfam; PF00007; Cys_knot; 1.
                  Ephydroidea; Drosophilidae; Drosophila
NCBI_TaxID=7224;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR006208; Cys_knot.
Pfam; PF00007; Cys_knot; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-JUN-1998 (TrEMBLrel. 06, Created)
01-JUN-1998 (TrEMBLrel. 06, Last sequence update)
01-JUN-2093 (TrEMBLrel. 24, Last samotation update)
Luteinizing hormone/chorionic gonadotrophin beta-subunit
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4332 MW;
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Laurasiatheria; Perissodactyla;
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Last
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EMBL/GenBank/DDBJ
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Pred.
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Pred. No. 2.2e+0
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ches 0;
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                                                                                     Insecta;
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                                                                   Muscomorpha;
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                                                                                        Pterygota;
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Equidae; Equu
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Q23948 DACHY
ID Q23948;
DT 01-NOV-1
DT 01-NOV-1
DT 01-MAR-2
DE Testis-s-
GN Name=Mst
OS Drosophi
OC Eukaryot
OC Neopterz
OC Ephydroi
OC MCBI_Ta
RN (1)
RP NUCLEOTI
RA Kuech A.
RL Submitte
DR F1yBase
SQ SEQUENCI
RESULT 7

QBMLN5 DROME
ID QBMLN5;
AC QBMLN5;
DT 01-OCT-2
DT 01-OCT-2
DT 13-SEP-2
DT 13-SEP-2
DT 13-SEP-2
DT 13-SEP-2
CG30430-
GN Name=CG3
OS Drosophi
OC Eukaryotto
OC Neoptera
OC Sphydroi
OX NCBI_Tax
RN [1]
RP NUCLEOTI
RX MEDLINE=
RA Adams M.
RA Amanatid
RA George R
RA George R
RA Sutton G
RA Ballaw R
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Matches 4
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Q23948;
Q13948;
Q1-NOV-1996 (TremBLrel. 01.
Q1-NOV-1996 (TremBLrel. 01.
Q1-MAR-2004 (TremBLrel. 26.
                     NUCLEOTIDE SEQUENCE.

                                                                                                                                                                                                                                                                                              Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilidae; Drosophila. NCBI TaxID=7227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kuech A., Buenemann H.;
Submitted (DEC-1990) to the EMBL/GenBank/DDBJ
EMBL; X58114; CAA41118 l; -; Genomic_DNA.
FlyBase; FBgn0015097; Dhyd\Mst87Fb.
SEQUENCE 49 AA; 4550 MW; 24758FEC51369F1D
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Kuech A., Buenemann H.;
Kuech A., Buenemann H.;
Submitted (DEC-1990) to the EMBL/GenBank/DDBJ databases.
EMBL; X58114; CAA41117.1; -; Genomic_DNA.
FlyBase; FBgn0015096; Dhyd\Mst87Fa.
FlyBase; FBgn0015096; Dhyd\Mst87Fa.
SEQUENCE 49 AA; 4610 MW; BE158FEC51369F01 CRC64;
                                                                                                                                                                                                                                                                                                                                                                               Name=CG30430; ORFNames=CG30430;
Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-OCT-2002 (TrEMBLrel. 22,
01-OCT-2002 (TrEMBLrel. 22,
13-SEP-2005 (TrEMBLrel. 31,
CG30430-PA (IP04475p).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name=Mst87Fb; Synonyms=DhTc2;
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NCBI_TaxID=7224;
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RA Dodson K., Doup L.B. Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Durbin K.J., Evangelista C.C., Ferrae C., Ferriera S., Fleischmann W.,
RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,
RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,
RA McIntosh G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA McIntosh S.M., Mwy M., Murphy B., Murphy L., Murny D.M., Nelson D.L.,
RA Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T.,
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Syirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
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Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.O.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
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Kaminker J.S., Bergman C.M., Kronmiller
Patel S., Frise E., Wheeler D.A., Lewis
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rtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Ch
erry J.M., Cawley S., Dahlke C., Davenport L.B., Davies B.
Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz
dson K., Doup L.B., Downes M., Dugan-Rocha S., Dunkov B.C.
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R., Stapleton M., Pacleb J.
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EMBL; AY549956; AAS55565.1; -; Genomic_DNA.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0042742; P:defense response to bacteria;
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(Rel. 26, Created)
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ic sperm protein Mst84Dc.

update)

Name=Mst84Dc; ORFNames=CG17945;

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RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Mannatides P.G., Scherer S.E., Holt R.A., Evans C.A., Galle R.F.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Brandon R.C., Rogers Y.-H.C., Blazed R.G., Champe M., Pfeiffer B.D.,
RA Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,
RA Abril J.F., Agbbyani A., An H.-J., Andrews-Pfeinnkoch C., Baldwin D.,
RA Bellew R.M., Basu A., Baxendale J., Bayraktarglu L., Bessley E.M.,
Ra Beson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Gherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Glorek K.C., Busen D.A., Butler H., Cadleu E., Center A., Chandra I.,
RA Gebalos B., Delchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Gebalos B., Delcher A., Domes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Golse C., Gabrielian A.E., Garg N.S., Gelbart W., Glasser K.,
RA Goldek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
Alali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Melson D.R., Nelson K., Nixon K., Nusskern D.R., Mechen H.,
RA Melson D.R., Nelson K., Nixon K., Musskern D.R., Smith T.,
RA Melson D.R., Nelson K., Stapleton M., Strong R., Sun B.,
Svirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,
RA Mang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun B.,
Spier E., Zhong F.N., Zhong W., Zhon W., Zhong S., Yao Q.A.,
RA Spier E., Janes G. R., Nixon K., Shang G., Zhoo Q., Zhong L.,
The genome sequence of Drosophila melanogaster.";
Schu H., Smith H.O.,
Spier D. Schull M., Schull M., Smith H.O.,
Spier D., Shand M., Schull M., Smith H.O.,
Spier D., Sh
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MEDLINE=92102953; PubMed=1684716; DOI=10.1016/0925-4773(91)90064-D;

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RA cluster of four genes selectively expressed in the male germ line of Drosophila melanogaster.";

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                                                                                                                                                                                                                                                                      MEDLINE=22426069; PubMed=12537572;
Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.
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Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
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Bukaryota; Metazoa; Arthropoda; Hexapoda; Inse
Neoptera; Endopterygota; Diptera; Brachycera;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                              systematic review
                                                                                                                                                                                                                                          Lewis S.E.;
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tione Biol. 3:RESEARCH0083.1-RES
TISSUE SPECIFICITY: Testis.
DEVELOPMENTAL STAGE: Primary
DOMAIN: This protein in
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InterPro; IPR001450; 4Fe4S ferredoxin.
InterPro; IPR001634; MSSP-
InterPro; IPR001007; VWF_C.
Pfam; PF03940; MSSP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Name=Mst84Dc; ORFNames=CG17945;
Drosophila simulans (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygo
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
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QGQAT9;
05-JUL-2004 (TrEMBLrel. 2:
05-JUL-2004 (TrEMBLrel. 2:
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Ensembl; CG17945; Drosophila n
FlyBase; FB900004174; Mst84Dc.
InterPro; IPR005634; MSSP.
Pfam; PF03940; MSSP; 1.
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PROSITE; PS00198; 4FE45 FERREDOXIN; UNKNOWN_1

PROSITE; PS01208; VMFC_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       simulansand D. mauritiana.";
J. Mol. Evol. 59:277-282(2004).
EMBL; AY549957; AAS55566.1; -;
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MST84DC.
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"Association of misexpression with
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                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PubMed=15486701; DOI=10.1007/s00239-004-2622-y;
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MEDLINE=20196006; PubMed=10731132; DOI=10.1126/science.287.5461.2185; Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D., Ananatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F., George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Sutton G.G., Wortman J.R., Yandell G., Nelmon C.R., Miklos G.L.G., Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G., Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P., Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
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P08175; Q9VFQ6;
01-AUG-1988 (Rel. 08, Created)
01-AUG-1988 (Rel. 08, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Male specific sperm protein Mst87F.
Male specific sperm protein Mst87F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kuhn R., Schaefer U., Schaefer M.;
"Cis-acting regions sufficient for spermatocyte-specific transcriptional and spermatid-specific translational cont Drosophila melanogaster gene mst(3)gl-9.";
EMBO J. 7:447-454(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=Canton-S;
MEDLINE=88211557; PubMed=2835228;
Kuhn R., Schaefer U., Schaefer M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Drosophila mauritiana (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Drosophila melanogaster (Fruit fiy).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ephydroidea; Drosophilidae; Drosophila
NCBI_TaxID=7227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROSITE; PS00198; 4FE4S_FERREDOXIN; UNKNOWN_1.
PROSITE; PS01208; VWFC_1; 1.
SEQUENCE 55 AA; 5165 MW; 95A12F27D07B6F6C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              J. MOL. Bvol. 59:277-282 (2004).
EMBL; AY549955; AAS55564.1; -; Genomic InterPro; IPR001450; 4Fe48 ferredoxin.
InterPro; IPR005634; MSSP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Association of misexpression with sterility in hybrids of Drosophila simulansand D. mauritiana.";
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                                                                                                                                                                                                                                                                                                                                                                           STRAIN=Berkeley;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Name=Mst87F; Synonyms=mst(3)gl-9; ORFNames=CG17956;
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_DROME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PubMed=15486701; DOI=10.1007/s00239-004-2622-y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Name=Mst84Dc;
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27,
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Last annotation updat
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Pred. No.
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RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.,
RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
RA Hastin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Jalali M., Malush R., Karpen G.H., Ke Z., Kulp D., Lai Z.,
Liu X., Mattei B., McIncosh T.C., McLeod M.P., McPherson D.,
RA Liu X., Mattei B., McIncosh T.C., McLeod M.P., McPherson D.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Spier E., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,
RA Spier E., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Wang Z.-Y., Wassarman D.A., Weinstcock G.M., Weissembach J.,
RA Yellon S.M., Woodseg T., Worley K.C., Wu D., Yang S., Yao Q.A.,
RA Yellos R.A., Myers E.W., Rubin G.M., Jang G., Zhao Q., Zheng L.,
RA Gibbs R.A., Myers E.W., Rubin G.M., Venter J.C.,
The genome sequence of Drosophila melanogaster.";
RT Trie genome sequence of Drosophila melanogaster.";
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                                                                                                  Query Match
Best Local (
                                                                               Matches
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-!- TISSUE SPECIFICITY: Testis.
-!- DEVELOPMENTAL STRACE: primary spermatocytes.
-!- DOMAIN: This protein is mostly composed of repe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=22426069; PubMed=12537572;
Milsra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E.
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B.
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.O.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                                                             Developmental protein; Differentiation; Multigene Spermatogenesis. SEQUENCE 56 AA; 5233 MW; 830CD13212C34A47 CRC6
                                                                                                                                                                                                                                                                    PIR; S00340; WTFF.
HSSP; P01180; 1JK6
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                                                                                                                                                                                                                         Ensembl; CG17956; Drosophila
FlyBase; FBgn0002862; Mst87F
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                                                                                                                                                                                                                                                                                                        Y00831; CAA68761.1; -; Genomic DNA.
AE003702; AAF54994.1; -; Genomic DNA
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Pred. No.
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EMBL outstation -
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RESULT 13 Q595V9_NEILA

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Q5F8A3_NEIG1
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A Lewis L.A., Gillaspy A.F., McLaughlin R.E., Gipson M., Ducey T.F.,

A Cownbey T., Hartman K., Nydick C., Carson M.B., Vaughn J., Thomson C.,

A Song L., Lin S., Yuan X., Najar P., Zhan M., Ren Q., Zhu H., Qi S.,

A Kenton S.M., Lai H., White J.D., Clifton S., Roe B.A., Dyer D.W.;

The complete genome sequence of Neisseria gonorrhoeae.";

Submitted (MAR-2003) to the EMBL/GenBank/DDBJ databases.

EMBL; AE004969; AAW89584 1; -; Genomic DNA.

GO; GO:0005489; F:metal ion binding; IEA.

GO; GO:0006118; P:metal ion binding; IEA.

GO; GO:0006118; P:electron transporter activity; IEA.

GO; GO:0006118; P:electron transport; IEA.

R GO; GO:0010152; Rubredoxin.

R InterPro; IPR0010152; Rubredoxin.

Pfam; PF00301; Rubredoxin.
Query Match
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10-MAY-2005
10-MAY-2005
10-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the neisserial gene pool.";

Submitted (JAN-2004) to the EMBL/GenBank/DDBJ databases.

EMBL; AY532628; AAW21818.1; -; Genomic DNA.

GO; GO:0005489; F:electron transporter activity; IEA.

GO; GO:0046872; F:metal ion binding; IEA.

GO; GO:0006118; P:electron transport; IEA.

Electron transport; Iron; Metal-binding; Rubredoxin.
                                                                                                                                                             PRINTS; PRO0163; RUBREDOXIN.
ProDom; PD001610; Rubredox; 1.
PROSITE; PS00202; RUBREDOXIN;
PROSITE; PS50903; RUBREDOXIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Q5F8A3_NEIG1
Q5F8A3;
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                                                                                            Rubredoxin.
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=242231;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Neisseria gonorrhoeae (strain ATCC 700825 / FA 1090).
Bacteria, Proteobacteria; Betaproteobacteria; Neisser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Identification of anomalous sequences in
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Bacteria; Proteobacteria;
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Similarity 4; Conser
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                                                                                                                                                                PS00202; RUBREDOXIN; 1.
PS50903; RUBREDOXIN_LIKE; 1.
                                                                                              56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           56 AA;
larity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bart A., Luyf A.C.M., van Kampen A.H.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6345 MW;
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                                                                                                                                        transport;
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                                                                                            17DACAA86E7DB8CF CRC64;
                        Score 31;
Pred. No.
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Pred. No. 3e+02;
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                        3e+02;
                                              DB 2;
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                                           Length 56;
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RESULT 16
Q9JQN3_NEIMA
ID Q9JQN3_NEIMA PRELIMINARY;
ID Q9JQN3;
AC Q9JQN3;
DT 01-OCT-2000 (TrEMBLrel. 15, L;
DT 01-OCT-2003 (TrEMBLrel. 25, L;
DT 01-OCT-2003 (TrEMBLrel. 25, L;
DE Putative rubredoxin.
GN OrderedLocusNames=NMA1201;
OS Neisseria meningitidis (serogion Neisseria; Beta OC Neisseriaceae; Neisseria.
OX NCBI_TaxID=65699;
                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tettelin H., Saunders N.J., Heidelberg J.F., Jeffries A.C., Nelson K.E., Eisen J.A., Ketchum K.A., Hood D.W., Peden J.F., Dodson R.J., Nelson W.C., Gwinn M.L., DeBoy R.T., Peterson J.D., Hickey E.K., Haft D.H., Salzberg S.L., White O., Fleischmann R.D., Dougherty B.A., Mason T.M., Ciecko A., Parksey D.S., Blair E., Cittone H., Clark B.B., Cotton M.D., Utterback T.R., Khouri H.M., Qin H., Vamathevan J.J., Gill J., Scarlato V., Mason E.R., Rappuoli R., Venter J.C.;
                                            Neisseria meningitidis (serogroup
                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR004039; Rubredox.
InterPro; IPR001052; Rubredoxin.
Pfam; PF00301; Rubredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GO; GO:0005489; F:electron transporter activity; IEA.
GO; GO:0046872; F:metal ion binding; IEA.
GO; GO:0006118; P:electron transport; IEA.
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Neisseria meningitidis (serogroup
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; AE002098;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Science 287:1809-1815(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=MC58 / Serogroup B;
MEDLINE=20175755; PubMed=10710307; DOI=10.1126/science.287.5459.1809;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                       PRINTS; PR00163; RUBREDOXIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Neisseriaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Complete genome sequence of Neisseria meningitidis serogroup B strain
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4 (TrEMBLrel. 27,
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ilarity 100.0%;
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RUBREDOXIN_LIKE; 1.
; Electron transport; Iron; Metal-binding;
                                                                                                                                                                                                                                                                                                                                                                                                           Rubredox; 1.
                                                                                                                                                                                                                                                                                                                                     6369
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                               Betaproteobacteria; Neisseriales;
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Last annotation updat
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Pred. No. 3e+
0; Mismatches
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                                                                                                                                                                                                                                                                                         3e+02;
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RESULT 17
QGQATG DROSI
ID OFQATG PROSI PRELIMINARY;
AC QGQATG;
DT 05-JUL-2004 (TrEMBLrel. 27,
DR 05-JUL-2004 (TrEMBLRel. 27,
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Matches 4
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STRAIN=Valparaiso;
STRAIN=Valparaiso;
STRAIN=Valparaiso;
STRAIN=Valparaiso;
STRAIN=Valparaiso;
STRAIN=Valparaiso;
Michalak P., Noor M.A.F.;
Michalak P., Noor M.A.F.;
"Association of misexpression with sterility in hybrids of Drosophila simulansand D. mauritiana.";
J. Mol. Evol. 59:277-282 (2004).
EMBL; AY549958; AAS55567.1; -; Genomic_DNA.
InterPro; IPR001450; 4Fe4S_ferredoxin.
InterPro; IPR001007; VWPC_T; I.
PROSITE; PS00198; 4FE4S_FERREDOXIN; UNKNOWN_1.
PROSITE; PS01208; VWPC_T; 1.
SEQUENCE 58 AA; 5422 MW; 8C70EDCC7E69AC83 CRC64;
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Whitehead S., Spratt B.G., Barrell B.G.;
"Complete DNA sequence of a serogroup A strain of Neisseria"
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NUCLEOTIDE SEQUENCE.
STRAIN=Z2491 / Serog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Name=Mst84Dc; ORFNames=CG17945;
Drosophila simulans (Fruit fl);
Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygo
Bukaryota; Endopterygota; Diptera; Brachycera; Muscomorpha;
Benydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PIR; H81133; H81133.
HSSP; P24297; 1RWD.
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EMBL; AL162755; CAB84462.1; -;
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MEDLINE=20222556; PubMed=10761919; DOI=10.
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PRODOM; PD001610; Rubredox; 1.
PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
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InterPro; IPR001052; Rubredoxin.
                                                                            Local Similarity
ses 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GO:0005489; F:electron transporter activity; GO:0046872; F:metal ion binding; IEA. GO:0006118; P:electron transport; IEA.
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4; Conserv
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Pred.
                                                                                                              Score 31; DB 2;
Pred. No. 3.1e+02
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RA Zeng
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RT Toxic
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                                                                                                                               Signal;
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EMBL; AF079059; AAD47373.1; -; Genomic_DNA.
EMBL; AF159976; AAK61823.1; -; mRNA.
EMBL; AF419252; AAN32699.1; -; Genomic_DNA.
HSSP; P15222; 1SIS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shunyi Z., Wenxin L.; "cDNA encoding a putative insect toxin from BmK."; Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE=Venom gland;
MEDLINE=20175769; PubMed=10708793; DOI=10.1016/S0041-0101(99)00212-3;
Zeng X.-C., Li W.-X., Zhu S.-Y., Peng F., Zhu Z.-H., Wu K.-L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mesobuthus martensii (Manchurian scorpion) (Buthus marte
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida;
Buthida; Buthoidea; Buthidae; Mesobuthus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Neurotoxin BmKCT precursor (C1-channel toxin)
chain toxin KCT) (BmKCL1) (TYCL1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SCCT_MESMA
Q9UADO;
                                                                                                                                                                                                                                                                                                                                           This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the El the European Bioinformatics Institute. There are no rest
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Submitted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE.
Wu J.-J., Dai L., Chi
Submitted (JUL-1998) t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=34649;
                              SEQUENCE
                                                                                                                                                               Chloride
                                                                                                                                                                              Pfam; PF05294; Toxin
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5294; Toxin_5; 1.
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                            Neurotoxin BmKCT.
By similarity.
By similarity.
By similarity.
By similarity.
C9FAABAFCB8E02EA C
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Length
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RESULT 20
Q71DA0_DROYA
ID Q71DA0_DROYA PRELIMINARY; P
AC Q71DA0_DROYA PRELIMINARY; P
AC Q71DA0;
DT 05-JUL-2004 (TrEMBLrel. 27, Crea
DT 05-JUL-2004 (TrEMBLrel. 27, Last
DT 05-JUL-2004 (TrEMBLrel. 27, Last
DT 05-JUL-2004 (TrEMBLrel. 27, Last
DE CG4959 (Fragment).
OS Drosophila yakuba (Fruit fly).
OC Eukaryota; Metazoa; Arthropoda; I
OC Neoptera; Endoptera; Diptera;
OC Ephydroidea; Drosophilidae; Droso
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SCCX_ME
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Matches 4; Conserv
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"A novel neurotoxin from scorpion BmK.";

Submitted (DEC-2000) to the EMBL/GenBank/DDBJ
-I- FUNCTION: Chloride channel ligand. Blocks
-chloride channels (By similarity).
-i- SUBCELLULAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DISULFID DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Signal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence up
10-MAY-2005 (Rel. 47, Last annotation
Neurotoxin Bml2-b precursor:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MESMA
  Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygo
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This Swiss-Prot entry is copyright. It is produced through a obstween the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
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Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Scorpi
Buthida; Buthoidea; Buthidae; Mesobuthus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q9BJW4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE SPECIFICITY: Expressed by the venom SIMILARITY: Belongs to the short scorpion (channel inhibitor subfamily.
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5294; Toxin_5; 1.
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Neurocoxin Bm12-b.
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Pred. No. 3.2
0; Mismatches
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RESULT 22

Q8T403 DROME
ID Q8T403; DROME PRELIMINARY; PRT; 61 AA.
AC Q8T403; DT 01-JUN-2002 (TrEMBLrel. 21, Created)
DT 01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
DT 10-MAY-2005 (TREMBLrel. 30, Last annotation update)
DT AT2355Lp (CG31740-PA).
GN ORFNames=CG31740;
OC Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pte.
OC Neoptera; Endopterygota; Diptera; Brachycera; Muscomorj
OC Ephydroidea; Drosophilidae; Drosophila.
OX NCBI_TaxID=7227;
RN [1]
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Matches 4
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QBMSH6;
01-OCT-2002 (1
01-OCT-2002 (1
01-MAR-2004 (1
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Drosophila melanogaster (Fruit fly).
Drosophila melanogaster (Fruit fly).
Eukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota;
Neoptera, Endopterygota, Diptera, Brachycera, Muscomorpha,
Ephydroidea, Drosophilidae, Drosophila.
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                                                                                                                                                                                                                                                                                               EMBL; AY118808; AAM50668.1; -
FlyBase; FBgn0028883; CG4959.
SEQUENCE 60 AA; 6117 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GH21762p.
                                                                                                                                                                                                                                                                                                                                     Submitted (JUN-2002) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=Berkeley;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=7227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "An evolutionary analysis of orphan
Genome Res. 13:2213-2219(2003).
EMBL; AF531969; AAQ09868.1; -; mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=22887302; PubMed=1
Domazet-Loso T., Tautz D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=7245;
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RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.B., Lewis S.E., Holt R.A., Evans C.A., Gocayne J.D., Ra Amanatides P.G., Scherer S.E., Holt R.A., Evans C.A., Gocayne J.D., Ra George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., Gebrar S.C., Lewis S.E., Richards S., Ashburner M., Henderson S.N., Ra Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Ra Bardon R.C., Rogers Y.-H.C., Blazed R.G., Champe M., Pfeiffer B.D., Ra Barndon R.C., Rogers Y.-H.C., Blazed R.G., Champe M., Pfeiffer B.D., Ra Barlew R.M., Basu A., An H.-J., Andrews-Pfannkoch C., Baldwin D., Belle W.R.M., Basu A., Barker E.G., Helt G., Nelson C.R., Wiklos G.L.G., Ra Ballew R.M., Baros P.V., Berman B.P., Bhandari D., Bolshakov S., Burtier H., Gadieu B., Center A., Chandra I., Ra Burtis K.C., Busam D.A., Butler H., Gadieu B., Center A., Chandra I., Ra Burtis K.C., Busam D.A., Butler H., Gadieu B., Center A., Dietz S.M., Dodson K., Doup L.E., Downes M., Dugan Rocha S., Dunkov B.C., Dunn P., Ra Pollos R., Geng F., Gorrell J.H., Gu Z., Guan P., Harris M., Ra Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Ra Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Alasko P., Lei Y., Levitsky A.A., Li J.H., Li Z., Liang Y., Lin X., Li X., Wattei B., McIntosh T.C., Weis M.-H., In Synam C., Ra Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z., Liang Y., Lin X., Li X., Wattei B., McIntosh T.C., McLeed M.P., McPherson D. L., Ra Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., McHeston D.H., Ra Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Mchert K., Remington K., Saunders R.D. C., Scheler F., Shen H., Ra Spier E., Spradling A.C., Stapleton M., Strong R., Sun S., Shen H., Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J., Ra Spier E., Spradling A.C., Stapleton M., Strong R., Sin B., Yao Q.A., Ra Zheng X.H., Zhong F.N., Zhong W., Zhang M., Zhang G., Zhao Q., Zheng L., The Golden G., Sheng L., Shen H., Shen H.O., Shen H., Shen H., Shen H., Shen H., Shen H., Shen H., Shen H.
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Genome Biol. 3:RESEARCH0079-RESEARCH0079(2002).
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Patel S., Frise E., Wheeler D.A.,
Ashburner M., Celniker S.E.;
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  MEDLINE=22426069;
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                                                                                                        melanogaster
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P81761; Q8T0X4;
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EMBL; AY089413; AAL99151.1; -; mRNA.
EMBL; A2003657; AAN11177.1; -; Genomic_DNA
Ensembl; CG31740; Drosophila melanogaster.
FlyBase; FBgn0051740; CG31740.
SEQUENCE 61 AA; 5965 MW; 94390FEB1D8B4
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Celniker S., Carlson of
Hoskins R., Stapleton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S. Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A., Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q., Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lepidopteran-selective toxin precursor (ButaIT) (BTChl2).
Mesobuthus tamulus (Eastern Indian scorpion) (Buthus tamu
Eukaryota, Metazoa, Arthropoda, Chelicerata, Arachnida, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yu C., Rubin G.; "Drosophila melanogaster Submitted (MAR-2000) to t
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10-OCT-2003 (Rel. 42, Last sequence up
10-WAY-2005 (Rel. 47, Last annotation
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Newton K.A., Armugam
"Insect toxin from Me
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Buthida; Buthoidea;
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                                                                                                                                                                                                                                  Wudayagiri R.,
                                                                                                                                                                                                                                                                                                                                                       Submitted
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                                                                                                                                                         tamulus.";
                                                                                                                                                                                              "Isolation
                                                                                                                                                                                                                                                         PubMed=11782289;
                                                                                                                                                                                                                                                                           FISSUE=Venom;
                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
                                                                                         FUNCTION: Induces flaccid petoxic to S.falculata larvae
                                  SUBCELLULAR LOCATION: Secreted TISSUE SPECIFICITY: Expressed | MASS SPECTROMETRY: MW=3856.7; |
                                                                                                                                Biochem. 2:16-16(2001)
                      NOTE=Ref
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                                                                                                                                                                         characterization of a venom of South Indian
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on J., Wa
con M., P
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Wan K., Pfeiffer
, Pacleb J., Park
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EMBL/GenBank/DDBJ
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                                                                                                                  paralysis
                                      by the venom gland. METHOD=Electrospray;
                                                                                                  mice
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scorpion
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scorpion, Mesobuthus
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                                                                                                                    H.virescens
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Svirskas
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                                          RANGE=25-61;
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R., Smith
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A Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,

A Adams M.D., Celniker S.E., Li P.W., Hoskins R.A., Galle R.F.,

A George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,

Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,

A Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,

A Wan K.H., Doyle C., Baxter B.G., Helt G., Nelson C.R., Miklos G.L.G.,

RA Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D.,

RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,

RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,

RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,

RA Borkova D., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,

RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

RA de Pablos B., Delcher A., Dong Z., Mays A.D., Dew I., Dietz S.M.,

RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
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Matches 4
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A DROME STANDARD; PRT; 63 AA
Q01642; Q9VIA3;
01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                                                                         Male specific sperm protein Mst84Da.

Name=Mst84Da; ORFNames=CG17946;

Drosophila melanogaster (Fruit fly).

Eukaryota; Metazoa; Arrhropoda; Hexapoda; Insecta; Pterygota;

Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
                                                                                                                                                                                                                   MEDLINE=20196006; PubMed=10731132; DOI=10.1126/science.287.5461.2185;
                                                                                                                                                                                                                                 STRAIN=Berkeley;
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NCBI_TaxID=7227;
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Pred. No. 3.3e+02;
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RESULT 25
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AC Q8MI19;
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                               MEDLINE=22426069, PubMed=12537572;
Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Misra S., Crosby M.A., Mungall C.J., Millburn G.H., Prochnik S.E.
Hradecky P., Huang Y., Kaminker S.E., Millburn G.H., Prochnik S.E.
Smith C.D., Tupy J.L., Whitfield B.J., Bayraktaroglu L., Berman B.
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; X67703; CAA47937.1; -; EMBL; AE003672; AAF54022.1; -
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DEVELOPMENTAL STAGE: Primar
DOMAIN: This protein is mos
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                                                                                                                                                                                                                                                                                                                                         protein; Differentiation; Multigene family;
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G., Zhao Q., Zheng L.,
S., Zhu X., Smith H.O.,
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Tayade C., Black G.P., Croy A.;

Tayade C., Black G.P., Croy A.;

Submitted (MAY-2004) to the EMBL/GenBank/DDBJ dat

EMBL; AY616676; AAT38876.1; -; mRNA.

GO; GO:0016020; C:membrane; IEA.

GO; GO:000883; F:growth factor activity; IEA.

GO; GO:0008283; P:cell proliferation; IEA.

GO; GO:0000874; P:regulation of cell cycle; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUB=Placental artery endothelium; Chung J.-Y., Tsoi S.C.M., Wen Y.-X., Magness R.R., Zheng J.; Submitted (AUG-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AF534638; AAN04109.1; -; mRNA.
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01-OCT-2002 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                  NON TER
                                                                                                                                                                                                                                                                                                                                         PROSITE; PS502,0,
Growth factor; Mitogen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; AF534638; AAN SMR; Q8MI19; 15-64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pecora; Bovidae;
NCBI_TaxID=9940;
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Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia;
                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR000072; PD_growth_iactor.
Pfam; PF00341; PDGF; 1.
ProDom; PD001629; PD_growth_factor; 1.
PROSITE; PS50278; PDGF_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Suina; Suidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vascular endothelial growth
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Pred. No. (
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             PRT;
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PERRABAN REPRESENTE DE LA REPRESENTA DE LA RESERVA DE LA RESERVA
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01-MAY-2000 (TrEMI
01-MAY-2000 (TrEMI
10-MAY-2005 (TrEMI
CG17376-PB, isofo:
ORFNames=CG17376;
                                                                                                                                                                                                                                                                                                                                     Celniker S.E., Wheeler D.A., Kronmiller B., Carlson J.W., Halp Patel S., Adams M., Champe M., Dugan S.P., Frise E., Hodgson A George R.A., Hoskins R.A., Laverty T., Wuzny D.M., Nelson C.R. Pacleb J.M., Park S., Pfeiffer B.D., Richards S., Sodergren E. Svirskas R., Tabor P.E., Wan K., Stapleton M., Sutton G.G., Ve Weinstock G., Scherer S.E., Myers E.W., Gibbs R.A., Rubin G.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Science
[2]
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ephydroidea; Dr
NCBI_TaxID=7227
                                                                                                        MEDLINE-22426070; PubMed=12537573; MEDLINE-22426070; PubMed=12537573; Kaminker J.S., Bergman C.M., Kronm Patel S., Frise E., Wheeler D.A.,
                                                                                                                                                                                                                                                          "Finishing a whole-genome shotgun: release 3 of the Drosophila melanogaster euchromatic genome sequence."; Genome Biol. 3:RESEARCH0079-RESEARCH0079(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=22426065; PubMed=12537568;
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                                               "The transposable elements of the Drosophila melanogaster
                                                                                Ashburner M.,
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                          perspective.
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(TrEMBLrel. 13, Last sequence update)
(TrEMBLrel. 30, Last annotation update)
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3:RESEARCH0084.1-RESEARCH0084.20(2002)
                                                                                      Celniker S.E.;
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Hoskins R.A.,
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., Rubin
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25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Hypothetical protein OSJNBa0016D04.29.
Name=CSJNBa0016D04.29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stapleton M., Čárlson J., Chavez C., Frise E., Park S., Wan K., Yu C., Rubin G.M., Celniker S., Submitted (FEB-204) to the EMBL/GenBank/DDBJ d EMBL; AR003615; AAF52440.1; -; Genomic DNA. EMBL; BT011549; AAS15685.1; -; mRNA. Ensembl; CG17376; Drosophila melanogaster. FlyBase; FBgn0042189; CG17376.

SEQUENCE 67 AA; 6871 MW; B9442F2C7B567C76 C
                                                                                                                                                                                                                                                                                                                         Oryza sativa (japonica cultivar-group).
Eukaryota; Viridiplantae; Streptophyta; Er
Spermatophyta; Magnoliophyta; Liliopsida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=22426069; PubMed=12537572;
Misra S., Crosby M.A., Mungall C.J., Milburn G.H., Prochnik S.E.
Hradecky P., Huang Y., Kaminker J.S., Milburn G.H., Prochnik S.E.
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B.
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                               Submitted (OCT-2002) to the EMBL; AP005800; BAD29328.1;
                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE.
Sasaki T., Matsumoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q6EPU8;
25-OCT-2004
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                                                                                                                                                                                                                                                                                                         Spermatophyta; Magnoliophyta;
Ehrhartoideae; Oryzeae; Oryza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=22426069; PubMe
                                                                                      Hypothetical
                                                                                                                                                                       "Oryza sativa nipponbare(GA3)
"Clone:OSJNBa0016D04.";
                                                                                                                                                                                                                                                                           NCBI_TaxID=39947;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            QGEPU8_ORYSA PRELIMINARY;
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the EMBL/GenBank/DDBJ
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, Pacleb J., Park
                                                                                                                                                                                                                   Katayose
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                                                               9FBA6730BE5730A6 CRC64;
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Score 31;
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  3.6e+02;
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a; Poales; Poaceae;
                    DB 2;
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Svirskas
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L., Berman B.P.,
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Best Local :
                   STRAIN=Oregon-R;
MEDLINE=92102953; Pu
Kuhn R., Kuhn C., Bo
"A cluster of four g
Kuhn R., Kuhn on the Ruhn R., Kuhn of Drosophila
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NON TER
SEQUENCE
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                                                                                                                                                                                   Male specific sperm protein Mst84Dd.

Name=Mst84Dd; ORFNames=CG17935;

Drosophila melanogameser (Fruit fly).

Eukaryota; Metazoa; Arthropoda; Hexapoda; Inse
Neoptera; Endopterygota; Diptera; Brachycera;

Ephydroidea; Drosophilidae; Drosophila.
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01-MAR-2004 (TrEMBLrel. 26,
ENSAMGP00000023827 (Fragment
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Q7PK85;
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The Anopheles gambiae Sequence Committee;
Submitted (APR-2004) to the EMBL/GenBank/DDBJ
-i- CAUTION: The sequence shown here is derive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The Anopheles gambiae Sequence Committee, "Anopheles gambiae re-annotation."; Submitted (APR-2002) to the EMBL/GenBank/
                                                                                                    DEVELOPMENTAL
                                                                                                                               NUCLEOTIDE
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Anophelinae; Anopheles.
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                                                                                                                                                                    NCBI_TaxID=7227;
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ANOGA
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                   PubMed=1684716; DOI=10.1016/0925-4773(91)90064-D;
Boersch D., Glaetzer K.H., Schaefer U., Schaefer
r genes selectively expressed in the male germ lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6483 MW;
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a; Nematocera; Culicoidea; Culicidae;
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35:143-151(1991)

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RA Adams M.P., Celliker S.E., Holt K.A., Woldyne J.D.,
RA Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Gocayne J.D.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Chen L.X.,
RA Wann K.H., Doyle C., Baxter E.G., Helt G., Welson C.R., Miklos G.L.G.,
RA Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D.,
RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brysstein P., Botshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brysstein P., Botshakov S.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Perriaz C., Ferriera S., Fleischmann W.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Hostin D., Houston K.A., Howland T.J., Hernandez J.R., Houck J.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Kimmel B.E., McIntosh T.C., McLeod M.P., McPherson D.,
RA Rainzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,
RA Rainert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Rainert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Rainer E., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,
RA Shue B.C., Siden-Kiamos I., Singson M., Skupski M.P., Smith T.,
RA Shue B.C., Sheng F.N., Welson D.,
RA Williams S.M., Woldage T., Welley J., Zhan Q., Zheng L.,
RA Shue S., La K., Welson D.,
RA King G., Zhao Q., Zhen
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S., Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E., Smith C.D., Tupy J.L., Whitfield E.J., Bayrakkaroglu L., Berman B.P. Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A., Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q., Stabeleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
  Ensembl;
FlyBase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu S., Gibbs R.A., Myers E.W., Rubin G.M., Venter J.C.; "The genome sequence of Drosophila melanogaster."; Science 287:2185-2195(2000).
                                        EMBL; X67703; CAA47940.1; -; EMBL; AE003672; AAF54026.1; PIR; S25775; S25775.
                                                                                                                                                                                                This Swiss-Prot entry is copyright. It is produced between the Swiss Institute of Bioinformatics and
                                                                                                                                                                                                                                                                                                                             Genome Biol. 3:RESEARCH0083.1-RESEARCH0083.22(;
-!- TISSUE SPECIFICITY: Testis.
-!- DEVELOPMENTAL STAGE: Primary spermatocytes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=22426069; PubMe Misra S., Crosby M.A.,
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CG17935; Drosophila melanogaster FBgn0004175; Mst84Dd.
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QSTSU4;
01-FEB-2005 (TIEMBLIG1. 29,
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01-FEB-2005 (TIEMBLIG1. 29)
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Q4V3L2
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Developmental protein; Differentiation; Multigene family; Repeat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (APR-2004) to the
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  melanogaster (Fruit fly).
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PubMed=14671304; DOI=10.1126/science.1088727;

Methe B.A., Nelson K.E., Eisen J.A., Paulsen I.T., Nelson W.C.,

Heidelberg J.F., Wu D., Wu M., Ward N.L., Beanan M.J., Dodson R.J.,

Madupu R., Brinkac L.M., Daugherty S.C., DeBoy R.T., Durkin A.S.,

Gwinn M.L., Kolonay J.F., Sullivan S.A., Haft D.H., Selengut J.,

Davidsen T.M., Zafar N., White O., Tran B., Romero C., Forberger H.

Weidman J.F., Khouri H.M., Feldblyum T.V., Utterback T.R.,

Van Aken S.E., Lovley D.R., Fraser C.M.;

"Genome of Geobacter sulfurreducens: metal reduction in subsurface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stapleton M., Carlson J., Chavez C., Frise E., George R. Park S., Wan K., Yu C., Celniker S.; Submitted (MAY-2005) to the EMBL/GenBank/DDBJ databases. EMBL; BT023344; AAY55766.1; -; mRNA.
SEQUENCE 72 AA; 6840 MW; 8CE37449ESF88ABA CRC64;
                                               Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Complete proteome; Hypothetical protein. SEQUENCE 72 AA; 8191 MW; F2E9B550E9D43BB9 CRC64;
    Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                          Name=Pkd1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TIGR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Science
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Geobacteraceae; G
NCBI_TaxID=35554;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Geobacter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUL-2004
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Ephydroidea; Drosophilidae; Drosophila.
                                                                                             Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               environments.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OrderedLocusNames=GSU1308;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GSU1308;
                                                                                                                                                                                                                                                                                                                                38
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AE017180; AAR34684.1; -; Genomic_DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10
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                                                                                                                G (TrEMBLrel. 31, CTREMBLrel. 31, CTREMBLrel. 31,
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                                                                                                                                                                                                           PRELIMINARY;
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Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Glires; Rodentia; Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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                                                                                                                Created)
Last sequence
Last annotation
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Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                 Score 31; DB 2;
Pred. No. 3.7e+02;
Mismatches 0;
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Pred. No.
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annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 72;
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RESULT 35
Q6EPU3 ORYSA PRELIMINARY; PRT; 73 AA.
ID Q6EPU3 ORYSA PRELIMINARY; PRT; 73 AA.
AC Q6EBU3;
DT 25-OCT-2004 (TrEMBLrel. 28, Created)
DT 25-OCT-2004 (TrEMBLrel. 28, Last sequence up
DT 25-OCT-2004 (TrEMBLrel. 28, Last annotation
DT 25-OCT-2004 (TrEMBLrel. 28, Last annotation
DT 25-OCT-2004 (TrEMBLRel. 28) Last annotation
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RY NUCLEOTIDE SEQUENCE.

RISSUB-Jaw and Limb;

RX MEDIINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

RX MEDIINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

RX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RX Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Wadin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Richards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Willalno D.K., Muany D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Query Match
Best Local S
Matches 4
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Best Local Similarity
Matches 4; Conserv
                                                                                                                Hypothetical SEQUENCE 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Name=OSJNBa0016D04.34;
Oryza sativa (japonica cultivar-group).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
                                                                                                                                                                      Gramene;
                                                                                                                                                                                             Submitted (OCT-2002) to the EMBL; AP005800; BAD29333.1;
                                                                                                                                                                                                                                                   clone:OSJNBa0016D04."
                                                                                                                                                                                                                                                                                                              Sasaki T., Matsumoto
                                                                                                                                                                                                                                                                                                                                                                                                                            Spermatophyta; Magnoliophyta; Liliopsida; Ehrhartoideae; Oryzeae; Oryza.
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NCBI TaxID=10090;
[1] ~.
                                                                                                                                                                                                                                                                                    Sasaki T., Matsumoto T., Katayose
"Oryza sativa nipponbare(GA3) geno
                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hypothetical protein.
SEQUENCE 72 AA; 7108 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IIISSUE=Jaw and Limb;
NIH MGC Project;
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Similarity 4; Conserv
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100.0%; ilarity 100.0%; Conservative 0;
                                                                                                          protein.
3 AA; 7746 MW;
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Score 31; DB
Pred. No. 3.8
0; Mismatches
                                                                                                                3EF220FB4F36D10A CRC64;
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Pred. No. 3.7e+02;
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                          DB 2; 1
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                                                     Length
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Best Local S
Matches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE-85122780; PhbMed=2578818; DOI=10.1016/0167-4781(85)90049-1; Wilton S.D., Crocker L.A., Rogers G.E.; Wilton S.D., Crocker L.A., Rogers G.E.; "Isolation and characterisation of keratin mRNA from the scale epidermis of the embryonic chick."; Biochim. Biophys. Acta 824:201-208(1985). EMBL; M25641; AAA48932.1; -; mRNA. PIR, A60136; A60136; A60136; GO:0005882; C:intermediate filament; IEA. GO; GO:0005882; C:intermediate filament; IEA. GO; GO:000580; F:structural constituent of cytoskeleton; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q9YH54 CHICK PRELIMINARY;
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01-JUN-2003 (TrEMBLrel.
Keratin (Fragment).
Gallus gallus (Chicken).
                                                                                                 STRAIN=Oregon-R;
MEDLINE=92102953; PubMed=1684716; DOI=10.1016/0925-4773(91)90064-D;
Kuhn R., Kuhn C., Boersch D., Glaetzer K.H., Schaefer U., Schaefer
"A cluster of four genes selectively expressed in the male germ lin
of Drosophila melanogaster.";
Mech. Dev. 35:143-151(1991).
                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE
DEVELOPMENTAL STAGE.
                                                                                                                                                                                                                                                                                                                                                       Neoptera; Endopterygota; Dip
Ephydroidea; Drosophilidae;
                                                                                                                                                                                                                                                                                                                                                                                                    Male specific sperm protein Mst84Db.

Name=Mst84Db; ORFNames=CG17934;

Drosophila melanogaster (Fruit fly).

Drosophila melanogaster (Fruit fly).

Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
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                                                    NUCLEOTIDE
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                             STRAIN=Berkeley
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                                                    SEQUENCE [LARGE SCALE GENOMIC DNA].
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PubMed=10731132; DOI=10.1126/science.287.5461.2185;
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Neognathae; Galliformes; Phasianidae; Phasiani
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Pred. No. 3.8e+02;
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RA Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G., Ra Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., Ra Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., Ra Baceson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Ra Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P., Ra Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P., Ra Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I., Ra Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I., Ra Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., Ra Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., Ra Glodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., Ra Glodsk A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Glasser K., Ra Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Ralush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A., Ra Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z., Liang Y., Lin X., Ra Liu X., Mattei B., McIntosh T.C., McLedd M.P., McPherson D., Ra Mcrkulov G., Milshina N.V., Mobarry C., Morris J., Pacleb J.M., Ra Rainert K., Remington K.A., Nixon K., Nusskern D.R., Pacleb J.M., Ra Rainert K., Remington K., Saunders R.D.C., Scheeler F., Spendling A.C., Stapleton M., Skupskin N.P., Smith T., Ra Shue B.C., Siden-Kiamos I., Simpson M., Skupskin N.P., Smith T., Ra Shue B.C., Siden-Kiamos I., Simpson M., Skupskin N.P., Smith T., Ra Shue B.C., Stapleton M., Strong R., Smith T., Ra Shue B.C., Stapleton M., Strong R., Smith T., Ra Shue B.C., Stapleton M., Strong R., Smith T., Ra Shue B.C., Stapleton M., Strong R., Smith T., Ra Shue B.C., Stapleton M., Strong R., Smith T., Ra Shue, R., Weissarman D.A., Weissenbach J., Wang S., Yao Q.A., Fan M., Jahang G., Zhao Q., Zheng L., Ra Khillians S.M., Woodage T., Wontley K.C., Mu D., Yang S., Yao Q.A., Shue G., Shue S., Shue S
                                                                                                                                                                                                                                                                                                                                                       Stapleton M., Carlson J.W., Brokstein P., Yu C., C George R.A., Guarin H., Kronmiller B., Pacleb J.M. Rubin G.M., Celniker S.E.; Rubin G.M., Celniker S.E.; "A Drosophila full-length cDNA resource."; Genome Biol. 3:RESEARCH0080.1-RESEARCH0080.8(2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=22426069; PubMed=12537572;
Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Misra S., Crosby M.A., Mungall C.J., Milburn G.H., Prochnik S.E.,
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B.
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=Berkeley;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genome Biol.
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Ref.4 sequence
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the use This between European as long a Swiss-Prot entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the Eleuropean Bioinformatics Institute. There are no rest as content is in no way modified and this statement restrictions EMBL ρı collaboration outstation 18

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RESULT 39
Q82DW2_STRAW ID Q82DW2_STRAW IAC Q82DW2;
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Best Local S
Matches 4
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Proc. Natl. Acad. Sci. U.S.A. 101:14925-14930(2004).

REMBL; APO06618; BAD57292.1; -; Genomic_DNA.

RGO; GO:0005489; F:electron transporter activity; IEA.

RGO; GO:0005506; F:iron ion binding; IEA.

RGO; GO:0006872; F:metal ion binding; IEA.

RGO; GO:0006118; p:electron transport; IEA.

RGO; GO:0006118; p:electron transport; IEA.

RINETRO, IPRO01450; 4Fe45 Fe_Sbd.

R InterPro, IPRO01450; 4Fe45 Fe_Sbd.

R PRINTS; PR00353; 4FE458FRDOXIN.

R PROSITE; PR00353; 4FE458FRDOXIN; 1.

R PROSITE; PS00198; 4FE45 FERREDOXIN; 1.

R PROSITE; PS00198; 4FE45 FERREDOXIN; 1.

M Metal-binding; Transport.

SEQUENCE 75 AA; 8193 MW; F6994795E7B2D209 CRC64;
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Best Local :
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PIR, S25773; S25773.
HSSP; OS9783; IEHH.
Ensembl; CG17934; Drosophila mel
FlyBase; FBgn0004173; Mst84Db.
InterPro, IPR005634; MSSP.
Pfam; PF03940; MSSP; 1.
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SEQUENCE
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STRAIN-IFM 10152;
PubMed=15466710; DOI=10.1073/pnas.0406410101;
Ishikawa J., Yamashita A., Mikami Y., Hoshino Shiba T., Hattori M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Spermatogenesis.
COMPBIAS 2
CONFLICT 42
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Q5YWZ9;
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Bacteria; Actinobacteria;
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25-OCT-2004 (TrEMBLrel.
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AE003672; AAF54024.1; -; Genomic DNA
AY113221; AAM29226.1; ALT_SEQ; mRNA.
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Pred. No. 3.9
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C -> W (in Ref. 1).
P -> L (in Ref. 1).
CC64F0F7A7E7CABC CRC64;
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Pred. No. 3.8e+02;
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RESULT 40

QARP51 TETING
QARP51 TETING PRELIMINARY; PRT; 75 AA.

ID QARP51 TETING PRELIMINARY; PRT; 75 AA.

AC QARP51;
AC QARP51;
DT 13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
DT 13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
DT 13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
DT 13-SEP-2005 (TremBlrel. 31
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NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

STRAIN=MA-4680 / ATCC 31267 / NCIMB 12804 / NRRL 8165;

STRAIN=21477403; PubMed=11572948; DOI=10.1073/pnas.211433198

OMNIAS S., Ikeda H., Ishikawa J., Hanamoto A., Takahashi C.,

Shinose M., Takahashi Y., Horikawa H., Nakazawa H., Osonoe T.

Kikuchi H., Shiba T., Sakaki Y., Hattori M.;

"Genome sequence of an industrial microorganism Streptomyces avermitilis: deducing the ability of producing secondary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRINTS; PRO0353; 4FE4SFRDOXIN.
PROSITE; PS00198; 4FE4S; FEREDOXIN;
4Fe-4S; Complete proteome; Electron
Metal-binding; Transport.
SEQUENCE 75 AA; 8139 MW; 8070930
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STRAIN-MA-4680 / ARTC 31267 / NCIMB 12804 / NRRL 8165;
STRAIN-MA-4680 / ARTC 31267 / NCIMB 12804 / NRRL 8165;
MEDILINE-22808306; PubMed=12692562; DOI=10.1038/nbt820;
Ikeda H., Ishikawa J., Hanamoto A., Shinose M., Kikuch Sakaki Y., Hattori M., Omura S.;
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01-JUN-2003 (TrEMBLrel. 24, Last sequ
01-MAR-2004 (TrEMBLrel. 26, Last anno
Putative ferredoxin.
Name=fdxE; OrderedLocusNames=SAV4856;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GO; GO:0005489; F:electron transporter activity; GO; GO:0005506; F:iron ion binding; IEA. GO; GO:0046872; F:metal ion binding; IEA. GO; GO:0006118; P:electron transport; IEA. InterPro; IPR001450; 4Fe4S_Pe_S_bd. Pfam; PF00037; Fer4; 2.
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Pred. No.
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transport; Iron;
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/pnas.211433198;
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RESULT
OF CONTROL

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Best Local S
Matches 4
                                                                                                                            Q74114_MYCPA PRELIMINARY;
Q74114;
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Summer E.J., Peek M.L., Haliburton J.R., Hall E., Simser J., No E.G., Gonzalez C.F., Young R.F.; Submitted (JUL-2003) to the EMBL/GenBank/DDBJ data EMBL; AY357582; AAQ63363.L; -; Genomic DNA. GO; GO:0019667; P.viral assembly, maturation, egre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
Putative head-tail joining protein.
Burkholderia cepacia phage BccpNazgul.
Viruses; dsDNA viruses, no RNA stage; Caudovirales
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-!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DBBJ whole genome shorgun (WGS) entry whi preliminary data.
EMBL; CAAE01015008; CAG09831.1; -; Genomic DNA.
                                                                                  Hypothetical protein.
OrderedLocusNames=MAP1106;
                                  Mycobacterium paratuberculosis.
Bacteria; Actinobacteridae;
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            Corynebacterineae;
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ProDom; PD019366; gpW; 1.
SEQUENCE 76 AA; 8345 MW;
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ilarity 100.0%;
Conservative (
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7913 MW; 4918F75E5D548359 C
            Mycobacteriaceae;
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                                  Actinomycetales;
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RESULT 44
Q4SJ47_TETNG I
ID Q4SJ47_TETNG I
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QOXHP3

ID WHP3

AC Q08

AC Q0
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01-MAR-2002 (TrEMBLrel. 20, 1

01-MAR-2002 (TrEMBLrel. 20, 1

01-JUN-2003 (TrEMBLrel. 24, 1

Probable glutaredoxin.

OrderedLocusNames=CPE2439;
                                                                                                                                                                                                                                                                                                                             Complete
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=1502;
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77 AA;
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            PRELIMINARY;
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flesh-eater.";

Proc. Natl. Acad. Sci. U.S.A. 99:996-1001(2002).

EMBL; BA000016; BAB82145.1; -; Genomic DNA.

GO; GO:0005489; F:electron transporter activity;

GO; GO:0006118; P:electron transport; IEA.
                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
STRAIN-13 / Type A;
MEDLINE-21664373; PubMed-11792842; DOI-10.1073/pnas.022493799;
Shimizu T., Ohtani K., Hirakawa H., Ohshima K., Yamashita A.,
Shiba T., Ogasawara N., Hattori M., Kuhara S., Hayashi H.;
"Complete genome sequence of Clostridium perfringens, an anaer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HSSP; P00214; IBOT.

GO; GO:0005489; F:electron transporter activity;
GO; GO:0005506; F:iron ion binding; IEA.
GO; GO:0046872; F:metal ion binding; IEA.
GO; GO:0046618; P:electron transport; IEA.
InterPro; IPR001450; 4Fe4S_Fe_S_bd.
Pfam; PF00037; Fer4; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae; Clostridium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRINTS; PR00353; 4FE4SFRDOXIN.
PROSITE; PS00198; 4FE4S FERREDOXIN;
4Fe-4S; Complete proteome; Electron
Metal-binding; Transport.
SEQUENCE 76 AA; 8289 MW; 8CF5D78
                                                                                                                           InterPro; IPR011911; GlrX_YruB.
InterPro; IPR006663; Thioredox_dom2
TIGREAMs; TIGR02196; GlrX_YruB; 1.
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EMBL/GenBank/DDBJ
-; Genomic_DNA.
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Last annotation update)
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Score 31; DB
Pred. No. 3.9
); Mismatches
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    transport; Iron; Iron-sulfur;

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                    .9e+02;
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Matches 4
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Jaillon O., Aury J.M., Brunet F., Petit J.L., Stange-Thomann N.,
A Mauceli E., Bouneau L., Fischer C., Ozouf-Costaz C., Bernot A.,
A Micaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,
A Nicaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,
A Nicaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,
A Dasilva C., Salanoubat M., Levy M., Boudet N., Castellano S.,
A Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
A Anthouard C., Skalli Z., Cattolico L., Poulain J., De Berardinis V.,
A Biemont C., Skalli Z., Cattolico L., Poulain J., De Berardinis V.,
A Cruaud C., Duprat S., Brottier P., Coutanceau J.P., Gouzy J.,
Parra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,
A Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,
A Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Lindbrad F.S., Weissenbach J., Roest Crollius H.;
"Genome duplication in the teleost fish Tetraodon nigroviridis reveals
"The Cattal Control of the Colonial Control of the Cattal Control of the Cattal Control of the Cattal Cattal Control of the Cattal Cattal Control of the Cattal Cattal Cattal Control of the Cattal Cattal
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Q9VJM5;
01-MAY-2000
01-MAY-2000
01-MAR-2004
NUCLEOTIDE SEQUENCE.

Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D., Adams M.D., Celniker S.E., Holt R.A., Boskins R.A., Gocayne J.D., Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Henderson S.N., Secret R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Sutton G.G., Balace J., Blazej R.G., Champe M., Pfeiffer B.D., Wan K.H., Doyle C., Baxter B.G., Helt G., Nelson C.R., Miklos G.L.G., Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., Abril J., Bassley E.M.,
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13-SEP-2005
13-SEP-2005
13-SEP-2005
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Submitted
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13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update chromosome undetermined SCAF14576, whole genome softwames=GSTENG00017363001;
                                                                                                                                                                                                                                                                                                                                            Name=CG13245; ORFNames=CG13245;
Drosophila melanogaster (Fruit fly)
Eukaryota; Metazoa; Arthropoda; Hexapoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tetraodon nigroviridis (Green puffer).

Bukaryota; Metazoa; Chordata; Craniata; Verte
Actinopterygii; Neopterygii; Teleostei; Eutel
Acanthomorpha; Acanthopterygii; Percomorpha;
Tetradontoidea; Tetraodontidae; Tetraodon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL;
                                                                                                                                                                                                                                                                                                                             Neoptera;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               45
DROME
                                                                                                                                                                                                                                                                                                                                                                                                                                                   L-MAY-2000 (TrEMBLrel.
L-MAY-2000 (TrEMBLrel.
MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               oscope; Whitehead Institute Centre for Genome Research; nitted (FEB-2004) to the EMBL/GenBank/DDBJ databases. CAUTION: The sequence shown here is derived from an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           preliminary data
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NCE 77 AA;
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ea; Drosophilidae;
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9043 MW; 25F48C1751A7D3A3 CRC64;
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13,
26,
                                                                                                                                                                                                                                                                                                     Diptera; Brachycera; Muscomorphā;
ae; Drosophila.
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Last sequence update)
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Pred. No.
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Euteleostei;
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RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.,
RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Glodek A., Gong F., Gorrell J.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D.A., Heiman T.J., Wei M.-H., Ibegwam C.,
RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
Liang Y., Lin X.,
RA Lasko P., Lei Y., Levitsky A.A., Li J.H., Li Z., Liang Y., Lin X.,
RA Lasko P., Lei Y., Levitsky A.A., Li J.H., Li Z., Liang Y., Lin X.,
RA Mchulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Neshrefi A.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Shue B.C., Staden-Kiamos I., Simpson M., Skupski M.P., Smith T.,
RA Syiers E., Waradinia A.C., Stapleton M., Strong R., Sun E.,
RA Yelson D.R., Weissenbach J.,
RA Yelson S.M., Wonder T., Zhong W., Zhong G., Zhong L.,
RA Yelson S.M., Wonder T., Song W., Shang G., Zhao Q., Zheng L.,
RA Yelson S.M., Warshan D.R., Zhan M., Zhan S., Zhu X.,
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MEDLINE=22426069; Pubmed=12537572;
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Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E.,
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B.P.,
Bettencourt B.R., Celniker S.B., de Grey A.D.N.J., Drysdale R.A.,
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"Drosophila melanogaster release Submitted (MAR-2000) to the EMBL/[6]
                                                                                  Berkeley Drosophila Genome Project;
Celniker S., Carlson J., Wan K., Pfeiffer
Hoskins R., Stapleton M., Pacleb J., Park
                                                                                                                                                                                                                 "Annotation of the systematic review."
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2426065; PubMed=12537568;
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Q95JG5 BOVIN
ID Q95JG5;
DT 01-DEC;
DT 01-DEC
RESULT 47
095JG6 BOVIN
ID 095JG6;
AC 095JG6;
DT 01-DEC-2001 (TrEMBLrel. 19,
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        expressed in cattle.";

Vet. Immunol. Immunopathol. 80:339-344(2001).

EMBL; AF322885; AAX73810.1; -; mRNA.

HSSP; O54709; 1HQ8.

SMR; Q95JG5; 1-80.
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Mammalia; Eutheria;
Pecora; Bovidae; Bov
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Submitted (MAR-2005) to
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MEDLINE=21351526; PubMed=11457486; DOI=10.1016/S0165-2427(01)00295-1;
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Q9VVA1:CG9706; NbExp=1; IntAct=EBI-85165,
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Sp. 9550041; C_TYPE_LECTIN_2;
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RESULT 48
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X MEDLINE=95283549; PubMed=7763258;

X MEDLINE=95283549; PubMed=7763258;

X MEDLINE=95283549; PubMed=7763258;

X Tahang F.P., Ramnikko A., Huhtaniemi I.;

I Usolation and characterization of testis-specific cDNAs for "Isolation and characterization" in the rate."

EMBL; U25803; AAC52251.1; -; mRNA.

R PIR; 165235; 165235.

R PIR; 165235; 165235.
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01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Testicular luteinizing hormone beta subunit.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pecora; Bovidae;
NCBI_TaxID=9913;
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01-OCT-2003 (TrEMBLrel. 25, 1
Natural killer cell receptor
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Mammalia; Eutheria;
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                                  InterPro; IPR006208; Cys_knot.
InterPro; IPR001545; Gly_hormc
Pfam; PF00007; Cys_knot; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROSITE; PS50041; C_TYPE_LECTIN_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TISSUE=Spleen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SM00034; CLECT;
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Euarchontoglires; Glires;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chordata; Crania
Laurasiatheria;
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Last annotation updat
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-; mRNA.
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Pred. No. 4.1
0; Mismatches
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lires; Rodentia; Sciurogna
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Euteleostomi;
Rhinocerotidae;

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RESULT
Q4T787
ID
TT787
AC
Q4T7187
AC
Q4T1113
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Q4T113
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Q4T1113
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ID 04
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DT 01
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Matches 4; Conser
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Best Local
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01-JUN-1998
01-JUN-2003
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B7 TETNG
Q4T7B7 TETNG
Q4T7B7;
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NON TER
NON TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tetraodon nigroviridis (Green puffer).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Actinopterygii; Meopterygii; Teleostei; Buteleostei; Neoteleostei;

Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;

Tetradontoidea; Tetraodontidae; Tetraodon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genoscope; Whitehead Institute Centre for Genome Research; Submitted (FEB-2004) to the EMBL/CenBank/DDBJ databases.
-i- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry whi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-SEP-2005 (TYEMBLrel. 31, Created)
13-SEP-2005 (TYEMBLrel. 31, Last sequence update)
13-SEP-2005 (TYEMBLrel. 31, Last annotation update)
Chromosome undetermined SCAP8161, whole genome sh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORFNames=GSTENG00005813001;
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SEQUENCE 80 AA; 8515 MW; F9EA66C2FD6FC97D CRC64;
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                                                                                                                                                                         _CERSI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CAUTION: The sequence shown EMBL/GenBank/DDBJ whole gence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   preliminary data.
, CAAE01008161; CAF91215.1;
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                                                                                                                                               CERSI
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4; Conserv
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80 AA;
                               (TrEMBLrel.
(TrEMBLrel.
(TrEMBLrel.
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                                                                                                                                               PRELIMINARY;
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8611 MW;
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Pred. No. 4.1
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        64705ABEA4A5620E CRC64;
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Pred. No.
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                         RESULT 52

O60744 HUMAN

O60744;

AC

O60744;

AC

O60744;

AC

O60744;

DT

O1-AUG-1998 (TrEMBLrel. 07, Cr

DT

O1-AWR-2004 (TrEMBLrel. 26, La

DT

O1-MAR-2004 (TrEMBLrel. 26, La

DE

Thioredoxin delta 3 (Fragment)

GN

Name=TXN delta 3;

OS

Homo sapiens (Human)

CE

Eukaryota; Metazoa; Chordata;
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Q42C47
Q472C.
AC Q472C.
AC
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Best Local S
Matches 4
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13-SEP-2005
13-SEP-2005
13-SEP-2005
Name=TXN delta 3;
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                             Thioredoxin delta 3 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dacteriophages.";

Proc. Natl. Acad. Sci. U.S.A. 102:5174-5179(2005)

EMBL, AY954959, AAX91390.1; -; Genomic DNA.

SEQUENCE 83 AA; 8546 MW; 710C2C35BBAB2C31 CRC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PubMed=15788529; DOI=10.1073/pnas.0501140102; Kwan T., Liu J., Dubow M., Gros P., Pelletier J.; "The complete genomes and proteomes of 27 Staphylococcus aureus
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NON_TER 1 1
SEQUENCE 82 AA; 8643 MW; 270411BA19
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Mammalia; Eutheria;
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Ceratotherium simum
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InterPro; IPR001545; Gly_hormoneB.
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to the EMBL/GenBank/DDBJ databases.
365.1; -; Genomic_DNA.
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neria; Perissodactyla;
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B1358B12.4 protein.
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SEQUENCE
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GO; GO:0005489; F:electron transporter activity; IEA.

GO; GO:0006118; P:electron transport; IEA.

InterPro; IPR006662; Thiored.

InterPro; IPR006663; Thioredox dom2.

InterPro; IPR012336; Thioredoxin-1ike.

InterPro; IPR012335; Thioredoxin, 1.

PERNITS; PR00421; THIOREDOXIN.
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SMR; C
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"The Genome of phiAsp2, an Actinoplanes Infecting Phage.";
Virus Genes 29:117-129(2004).
EMBL; AY576796; AAT36758.1; -; Genomic DNA.
SEQUENCE 84 AA; 8995 MW; CIECF1218IEB7C72 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.
NUCLEOTIDE SEQUENCE.
PubMed=15215690; DOI=10.1023/B:VIRU.0000032795.17713.6c;
Parling M., Bartkowiak K., Pape H., Meinhardt F.;
Parling M., Bartkowiak K., Pape H., Meinhardt F.;
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"An alternative splice variant of human
Chin. Sci. Bull. 43:292-295 (1998).
EMBL; AF065241; AAC17430.1; -; mRNA.
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05-JUL-2004
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Pred. No. 4.2e+02;
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Liu Y.L., Mu J., Yu Z., Chen L., Fan D.L., Weng Q.J., Zhang L.,
Lu Y.Q., Yu S.L., Liu X.H., Lu T.T., Zhang Y.J., Lu Y., Li C., Li T.,
Zhang Y., Hu H., Jia P.X., Qian Y.M., Ying K., Zhou B., Chen Z.H.,
Hao P., Zhang L., Wu M., Zhang R.Q., Guan J.P., Fu G., Wang S.Y.,
Ren S.X., Lu G., Lin W., Gu W.Q., Zhu G.F., Tu Y.F., Jia J., Yin H.F.,
Zhang Y., Cai Z., Chen J., Kang H., Chen X.Y., Shao C.Y., Sun Y.,
Hu Q.P., Zhang X.L., Zhang W., Wang L.J., Ding C.W., Sheng H.H.,
Gu J.L., Chen S.T., Ni L., Zhu F.H., Hong G.F.,
Gu J.L., Chen S.T., Ni L., Zhu F.H., Hong G.F.,
Submitted (NOV-2003) to the EMBL/GenBank/DDBJ databases.
EMBL, EX842605; CAE75995.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NÜCLEOTIDE SEQUENCE:

PubMed=12447439; DOI=10.1038/nature01183;

Peng Q. Zhang Y., Hao P., Wang S., Fu G., Huang Y., Li Y., Zhu J., Liu Y., Hu X., Jia P., Zhang Y., Zhao Q., Ying K., Yu S., Tang Y., Weng Q., Zhang L., Lu Y., Mu J., Lu Y., Zhang L.S., Yu Z., Fan D., Liu X., Lu T., Li C., Wu Y., Sun T., Lei H., Li T., Hu H., Quan J., Wu M., Zhang R., Zhou B., Chen Z., Chen L., Jin Z., Wang R., Yin H., Cai Z., Ren S., Lv G., Gu W., Zhu G., Tu Y., Jia J., Zhang Y., Chen J., Kang H., Chen X., Shao C., Sun Y., Hu Q., Zhang X., Zhang W., Mang L., Ding C., Sheng H., Gu J., Chen S., Ni L., Zhu F., Chen W., Lan L., Lai Y., Cheng Z., Gu M., Jiang J., Li J., Hong G., Xue Y., Li L., Lai Y., Cheng Z., Gu M., Jiang J., Li J., Hong G., Xue Y., Li L., Lai Y., Cheng Z., Gu M., Jiang J., Li J., Hong G., Xue Y.,
                         01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Type I hair keratin (Fragment).
Name=type I hair keratin;
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Han B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oryza sativa (japonica cultivar-group).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gramene; Q6MWE8; -.
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[1]
                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.
MEDLINE=95092656; PubMed=7528047;
                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                        Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR006461; DUF_A_thal_Cys.
Pfam; PF04749; PLAC8; 1.
SEQUENCE 84 AA; 9417 MW; F0BA5C5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                                                                                                                 Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                      Chordata; Craniata; Vertebra Euarchontoglires; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
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keratin
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                                                                                                                                                                                                                                                                                                                                                        Vertebrata; Euteleostomi; rimates; Catarrhini; Homin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.2e+02;
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                                                                                                                                                                            genes.";
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                                                                                                                                family.
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Gaps

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NUCLEOTIDE SEQUENCE.

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Best Local S
Matches 4
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Best Local S
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/10 9CAUD
QBW710 9CAUD 1
QBW710;
QBW710;
01-MAR-2002 (
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Q5T936;
01-FEB-2005 (
01-FEB-2005 (
01-FEB-2005 (
                                                                                                                                                                                                                                                                                         Cyanophage P60.
Viruses; dsDNA viruses,
NCBI_TaxID=151528;
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EMBL; AL158158; CAI14
SMR; Q5T936; 1-85.
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Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                  Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAR-2002
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Intermediate filament; Keratin.
NON TER 85
SEQÜENCE 85 AA; 9314 MW; 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GO; GO:0005489; F:electron transporter activity; GO; GO:0006118; P:electron transport; IEA. InterPro; IPR006662; Thiored. InterPro; IPR006663; Thioredox dom2. InterPro; IPR012336; Thioredox dom2. InterPro; IPR012336; ThioredoxIn-like. PF00085; Thioredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Williams S.
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-; Genomic_DNA.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              40.41B8DF0CC18414 CRC64;
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Pred.
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Primates; Cata
                                                                                                                                                                                                                                                                                                                                 Caudovirales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 85
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hes 0;
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                                 RESULT 59
THIO2 DICTO
ID THIO2
AC P2944
DT 01-AE
DT 10-MB
DE Thio2
GN Name-
OS Dicty
OC Eukal
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                                 Name=trxB; Synonyms=trx2;
Dictyostelium discoideum (Slime mold).
Eukaryota; Mycetozoa; Dictyosteliida;
                                                                                                                                    Thioredoxin
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RESULT
OPCENT
OP
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Maiti R., Ronning C.M
Barnstead M.R
Submit
                   D2 DICDI
THIO2 DICDI
P29446;
01-APR-1993
01-APR-1993
10-MAY-2005
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Q9C922;
01-JUN-2001 ()
01-JUN-2001 ()
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE.
Town C.D., Kaul S.;
Submitted (JAN-2001) to the
EMBL; AC019018; AAG52290.1;
HSSP; P23400; 1DBY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; AF338467; AAL73267.1; -; Genomic GO; GO:0005489; F:electron transporter GO; GO:0006118; P:electron transport; I Interpro; IPR006662; Thiored. PRINTS; PR00421; THIOREDOXIN.
SEQUENCE 87 AA; 9991 MW; FFBC315381
                                                                                                                                                                                                                                                                                                                                                                                                                      PRINTS; PR00421; THIOREDOXIN.
PROSITE; PS00194; THIOREDOXIN;
SEQUENCE 87 AA; 10318 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pfam;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GO; GO:0005489; F:electron transporter activity; GO; GO:0006118; P:electron transport; IEA. InterPro; IPR006662; Thiored. InterPro; IPR006663; Thioredox dom2. InterPro; IPR012336; ThioredoxIn-like. InterPro; IPR012335; ThioredoxIn-fold.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=3702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis thaliana (Mouse-ear cress).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence up
01-MAR-2004 (TrEMBLrel. 26, Last annotation
Putative thioredoxin; 109829-109566.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Name=F14G24.26;
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M.E., Bowman C.
(Rel. 25, Created)
(Rel. 25, Last sequence up
(Rel. 47, Last annotation
2 (Fragment).
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                                                                                                                 STANDARD;
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Pred. No. 4.4
); Mismatches
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Pred. No. 4.4
); Mismatches
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Dictyostelium.

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SETTITITIES REPRESENTATION OF THE PROPERTY OF 
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Best Local
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PRINTS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=92250653; PubMed=1577820; Wetterauer B., Jacquot J.-P., Veron M.; "Thioredoxins from Dictyostelium discoideum are a developmentally regulated multigene family.";
J. Biol. Chem. 267:9895-9904(1992).
-I- FUNCTION: Participates in various redox reactions through the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DISULFID NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Electron
ACT_SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DictyBase, DDB0001983; TRX2.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; M91382; AAA332
PIR; B46264; B46264.
HSSP; P10599; 1ERV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
NUCLEOTIDE SEQUENCE.
STRAINATCC 43049;
PubMed=15520287; DOI=10.1101/gr.2700304;
Baliga N.S., Bonneau R., Facciotti M.T.,
Deutsch E.W., Shannon P., Chiu Y., Weng R
Date S.V., Marcotte E., Hood L., Ng W.V.;
                                                                                                                                                                                                                                                                                                  01-FEB-2005 (TrEMBLrel 29,
01-FEB-2005 (TrEMBLrel 29,
01-FEB-2005 (TrEMBLrel 29,
                                                                                                                                                                                                                                                                                                                                                                        Q5V0J2 HALMA PRELIMINARY;
Q5V0J2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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                                                                                                                                                                                                                                  Name=trxA3; OrderedLocusNames=rrnAC2110;
Haloarcula marismortui (Halobacterium ma
                                                                                                                                                                                                                                                                                    Thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROSITE; PS00194; THIOREDOXIN;
                                                                                                                                                             NCBI_TaxID=2238;
                                                                                                                                                                                       Halobacteriaceae;
                                                                                                                                                                                                             Archaea;
                                                                                                                                                                                                                                                                                                                                                                                                                       HALMA
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FUNCTION: Participates in various redox reactions to reversible oxidation of its active center dithiol to and catalyzes dithiol-disulfide exchange reactions. SIMILARITY: Belongs to the thioredoxin family.

SIMILARITY: Contains 1 thioredoxin domain.
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eriaceae; Haloarcula.
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Nucleophile (By similarity).
Nucleophile (By similarity).
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ın R.R., H
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RESULT 61
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HSSP; P80579; INW2.
GO; GO:0005489; F:electron transporter a
GO; GO:0005418; P:electron transport; IE
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDXIN.
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Martinez-Arias R., Henne A., Wiezer A., Baeumer S.
Brueggemann H., Lienard T., Christmann A., Boemeck
Bhattacharyya A., Lykidis A., Overbeek R., Klenk H
Fritz H.-J., Gottschalk G.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=Goe1 / Go1 / ATCC BAA-199 / DSM 3647 / OCM MEDLINE=22120827; PubMed=12125824;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OrderedLocusNames=MM0436;
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E013269; AAM30132.1; -;
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, Boemecke M., Ste
, Klenk H.-P., Gun
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PubMed=15377793; DOI=10.1073/pnas.0403306101;

A Nierman W.C., DeShazer D., Kim H.S., Tettelin H., Nelson K.E.,

A Feldblyum T.V., Ulrich R.L., Ronning C.M., Brinkac L.M.,

A Daugherty S.C., Davidsen T.D., DeBoy R.T., Dimitrov G., Dodson R.A

A Durkin A.S., Gwinn M.L., Haft D.H., Khouri H.M., Kolonay J.F.,

A Madupu R., Mohammoud Y., Nelson W.C., Radune D., Romero C.M.,

A Sarria S., Selengut J., Shamblin C., Sullivan S.A., White O., Yu

A Sarria S., Selengut J., Shamblin C., Sullivan S.A., White O., Yu

7 "Structural flexibility in the Burkholderia mallei genome.";

Proc. Natl. Acad. Sci. U.S.A. 101:14246-14251(2004).

R EMBL; CP0000010; AAU50201.1; -; Genomic_DNA.
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GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0005179; F:hormone activity; IEA.
GO; GO:0005213; F:structural constituent of chorion
InterPro; IPR006208; Cys Knot.
InterPro; IPR001545; Gly_hormoneB.
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Pischer S., Veits J., Meyer H.H.D.;
Submitted (PEB-1998) to the EMBL/GenBank/DDBJ
Submitted (PEB-1998) to the EMBL/GenBank/DDBJ
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TOWNSHIP TEN
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SMART; SM00068; GHB; 1.

PROSITE; PS00689; GLYCO_HORMONE_BETA_2;
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25-OCT-2004 (TrEMBLrel.
25-OCT-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Name=LH/Cu-Deum,
Equus caballus (Horse).
Equus caballus (Horse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q71UV5_HORSE
Q71UV5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Burkholderiaceae;
NCBI_TaxID=13373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Burkholderia mallei (Pseudomonas mallei)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OrderedLocusNames=BMA2369;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hypothetical protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Luteinizing hormone/chorionic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (Fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        63
_BURMA
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HORSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   43
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BURMA PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 4; Conserv
   proteome;
88 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (TremBLrel. 27, Created)
(TremBLrel. 27, Last sequence update)
(TremBLrel. 27, Last annotation update)
hormone/chorionic gonadotrophin beta-subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Burkholderia.
   Hypothetical protein.
9445 MW; 74D511CDE7B2ECD6 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9153 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Laurasiatheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Betaproteobacteria; Burkholderiales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Created)
Last sequence update)
Last annotation updat
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Pred.
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heria; Perissodactyla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Nelson K.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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Equidae; Equus.
                                                                                                                                                                                                                                                                                                                                                   Dodson R.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       In.
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RESULT 65
046618 EQUZE
D46618 EQUZE PRELIMINARY;
AC 046618;
DT 01-JUN-1998 (TrEMBLrel. 06,
DT 01-JUN-1998 (TrEMBLrel. 24,
DT 01-JUN-2003 (TrEMBLrel. 24,
DE Luteinizing hormone/chorion
DE (Fragment).
GN Name=LH/GG-beta;
GN Name=LH/GG-beta;
CC Mammalia; Eutheria; Laurasi.
CC Mammalia; Eutheria; Laurasi.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 64
Q9HMD0 HALSA
ID Q9HMD0 H
AC Q9HMD0 T
D1 MAR-2
DT 01-MAR-2
DT 01-MAR-
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Best Local Similarity
Matches: 4; Conser
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Best Local
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Q9HMDO;
01-MAR-2001 (TrEMBLrel. 16,
01-MAR-2001 (TrEMBLrel. 16,
01-MAR-2004 (TrEMBLrel. 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=20504483; PubMed=11016950; DOI=10.1073/pnas.190337797; Ng W.V., Kennedy S.P., Mahairas G.G., Berquist B., Pan M., Shukla H.D., Lasky S.R., Baliga N.S., Thorsson V., Sbrogna J., Swartzell S., Weir D., Hall J., Dahl T.A., Welti R., Goo Y.A., Leithauser B., Keller K., Cruz R., Danson M.J., Hough D.W., Maddocks D.G., Jablonski P.E., Krebs M.P., Angevine C.M., Dale H., Isenbarger T.A., Peck R.F., Pohlschroder M., Spudich J.L., Jung K. Jahm M., Freitas T., Hou S., Daniels C.J., Dennis P.P., Omer A.D., Ebhardt H., Lowe T.M., Liang P., Riley M., Hood L., DasSarma S.; "Genome sequence of Halobacterium species NRC-1.", Proc. Natl. Acad. Sci. U.S.A. 97:12176-12181 (2000).
EMBL; AE005133; AAG20641.1; -; Genomic_DNA.
     Equus zebra hartmannae.
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Butheria; Laurasiatheria; Perissodactyla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PIR; E84409; E84409.
HSSP; P10599; 1ERV.
GO; GO:0005489; F:electron transporter activity;
GO; GO:0006118; P:electron transport; IEA.
GO; GO:0006810; P:transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
STRAIN-NRC-1 / ATCC 700922 / JCM 11081;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Halobacterium salinarium (Halobacterium halobium).
Archaea; Euryarchaeota; Halobacteria; Halobacteriales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRINTS; PR00421; THIOREDOXIN.
TIGRFAMs; TIGR01068; thioredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR006662; Thiored.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR006663; Thioredox_dom2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Name=trxA2; OrderedLocusNames=VNG2600G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Halobacteriaceae; Halobacterium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PF00085; Thioredoxin;
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1 (TrEMBLrel. 16,
4 (TrEMBLrel. 26,
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                                                                                                                                                hormone/chorionic
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                                                                                                                                          Last sequence update)
Last annotation updat
ic gonadotrophin beta-
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Last sequence update)
Last annotation updat
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Pred. No. 4.4
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 31; DB 2;
Pred. No. 4.4e+02;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BDF97B3496F70BFF CRC64;
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                                                                                                                                                update)
beta-subunit
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        Euteleostomi;
Equidae; Equu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>۰</u>
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RESULT
046619
RESULT 67
Q4HCJ2_9DEIO
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Best Local S
Matches 4
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Best Local
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PROSITE;
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NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                 SMART; SM
PROSITE;
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SEQUENCE
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GO; GO:0005179; F:hormone activity; IEA.
GO; GO:0005213; F:structural constituent
InterPro; IPR006208; Cys knot:
InterPro; IPR001545; Gly_hormoneB.
Pfam; PF00007; Cys knot; 1.
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Submitted (FBB-1998) to the EMBL/GenBank/DDBJ
EMBL; AF047602; AAC04360.1; -; Genomic_DNA.
HSSP; P01233; 1HCN.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0005179; F:hormone activity; IEA.
GO; GO:0005173; F:structural constituent of ch
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01-JUN-1998 (TYEMBLrel. 06, Last sequence update)
01-JUN-2003 (TYEMBLrel. 24, Last annotation update)
Luteinizing hormone/chorionic gonadotrophin beta-subunit
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01-JUN-1998
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Mammalia; Eutheria; Laurasiatheria; Perissodactyla;
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SMART; SM00068; GHB; 1.
                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR006208; Cys_knot.
InterPro; IPR001545; Gly_hormoneB
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GO:0005576; C:extracellular region; IGO:0005179; F:hormone activity; IEA.
GO:0005213; F:structural constituent
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EMBL/GenBank/DDBJ
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Q4LEN8;
13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
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Q4HCJ2;
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Name=10r-hep 2;

Paralichthys olivaceus (Japanese flounder).

Paralichthys olivaceus (Japanese flounder).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;

Acanthomorpha; Acanthopterygii; Percomorpha; Pleuronectiformes;

Pleuronectoidei; Paralichthyidae; Paralichthys.

NCBI_TaxID=8255;
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EMBL; AAHE01000001; EALB4178.1; -;
Hypothetical protein.
SEQUENCE 89 AA; 9776 MW; 1ABEA;
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STRAIN=DSM 11300;
US DOE Joint Genome Institute (JGI-ORNL);
Larimer F., Land M.;
"Annotation of the draft genome assembly of the Model of the Bank,"
Submitted (JUN-2005) to the EMBL/GenBank, I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hypothetical protein.
ORFNames=DgeoDRAFT_2284;
Deinococcus geothermalis DSM 11300.
Bacteria; Deinococcus-Thermus; Dein
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Coneland A., Lucas S., Lapidus A., Barry K.,
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STRAIN=DSM 11300;
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89 AA;
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98063; BAE06235.1; -; Genomic DNA.
89 AA; 9729 MW; 04CBB16152953F00
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                                         Conservative
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Pred. No. 4.4
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Best Local Similarity
Matches 4; Conserv
                                                                                                                              Paralichthys olivaceus (Japanese flounder).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Pleuronectiformes;
Pleuronectoidei; Paralichthyidae; Paralichthys.

NCBI TaxID=8255;
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Q4LEPO;
13-SEP-2005 (TrEMBLrel. 31, Created)
13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
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Giardia lamblia ATCC 50803.
Eukaryota; Diplomonadida; Hexamitidae; Giardiinae; Giardia.
NCBI_TaxID-184922;
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Submitted (JAN-2005) to the EMBL/GenBank/DDBJ databases.
EMBL; AB198061; BAE06233.1; -; mRNA.
SEQUENCE 89 AA; 9771 MW; D9AE7173C29CB412 CRC64;
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Hirono I., Nozaki R., Ono Y., Kurobe T., Ohira T., Aoki T.;
"Two different types of hepcidins from Japanese flounder Paralichthys
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25 80.6 1410 2 US-09-285-310-4 Sequenc 25 80.6 1410 2 US-09-753-385-4 Sequenc 25 80.6 1525 2 US-09-191-647-2 Sequenc 25 80.6 1525 2 US-09-540-245A-2 Sequenc	25 80.6 1404 2 US-09-195-524-2 Sequence 2, 25 80.6 1410 1 US-08-470-058-4 Sequence 4, 25 80.6 1410 2 US-09-037-188-4 Sequence 4,	25 80.6 1401 2 US-08-611-729A-2 Sequence 20 25 80.6 1404 1 US-08-611-729A-2 Sequence 2, 25 80.6 1404 2 US-08-611-729A-2	25 80.6 1276 4 PCT-US95-04228-24 Sequence 24 25 80.6 1347 2 US-09-058-489-24 Sequence 24	25 80.6 1276 2 US-08-446-648-24 Sequence 24 Sequence 24 Sequence 24	25 80.6 1276 1 US-08-222-616-24 Sequence 24	25 80.6 1257 1 US-08-340-428B-49 Sequence	25 80.6 1253 2 US-08-506-296B-14 Sequence 25 80.6 1253 2 US-09-489-039A-12097 Sequence	25 80.6 1240 2 US-09-058-489-23 Sequence 25 80.6 1240 2 US-09-058-489-23	25 80.6 1149 2 US-09-418-540-5 Sequence	25 80.6 1149 2 US-08-560-005-5 Sequence	25 80.6 1079 2 US-09-058-489-22 Sequence	25 80.6 1074 2 US-09-753-385-2 Sequence	25 80.6 1074 2 US-09-037-188-2 Sequence	25 80.6 1074 1 US-08-470-058-2 Sequence	25 80.6 1049 2 US-09-954-987B-172 Sequence 25 80.6 1049 2 US-10-020-445A-496 Sequence	25 80.6 1049 2 US-09-954-987B-170 Sequence	25 80.6 1024 2 US-09-949-016-7276 Sequence	25 80.6 991 2 US-09-949-016-7768 sequence 25 80.6 1024 2 US-09-562-737-88 Sequence	25 80.6 990 2 US-09-949-016-7235 Sequence	25 80.6 989 2 US-09-954-987B-171 Sequence	25 80.6 987 4 PCT-US95-08812-6 Sequence	25 80.6 987 1 US-08-436-044-6 Sequence	25 80.6 981 2 US-09-949-002-513 Sequence	25 80.6 970 1 US-08-673-789-7 Sequence	25 80.6 934 2 US-09-949-002-289 Sequence	25 80.6 863 2 US-09-619-353-14 Sequence	25 80.6 820 2 US-09-949-016-9224 Sequence	25 80.6 794 2 US-10-133-910-2 Sequence 2,	25 80.6 794 2 US-08-955-424-2 Sequence 2, 25 80.6 794 2 US-09-592-480-1 Sequence 1.	25 80.6 794 2 US-09-253-854-2 Sequence 2,	25 80.6 794 1 US-08-480-382-4 Sequence 4,	25 80.6 794 1 US-08-480-382-2 Sequence 2,	25 80.6 794 1 US-08-745-880-2 Seguence 2,	25 80.6 794 1 US-08-865-203-2 Sequence 2,	25 80.6 794 1 US-07-885-972A-2 Sequence 2,	25 80.6 771 2 US-09-188-930-183 Sequence 18	25 80.6 762 2 US-09-489-039A-10008 Sequence 10	25 80.6 729 2 US-09-252-991A-30945 Segmence 30	25 80.6 721 2 US-09-134-000C-5466 Sequence 54	25 80.6 719 2 US-08-753-247-12 Sequence	25 80.6 713 2 US-08-753-247-9 Sequence 9,	25 80.6 709 2 US-08-753-247-6 Sequence 6,

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RESULT 1
US-07-820-688A-5
              COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, Version #1.25

SOFTWARE: PATENTION DATA:

APPLICATION NUMBER: US/07/820,688A

FILING DATE: 19920324

CLASSIFICATION: 435

PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/NL91/00101

FILING DATE: 18-UTW-1991

PRIOR APPLICATION DATA:

APPLICATION NUMBER: EP 90201598.1

FILING DATE: 18-UTW-1990

ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: NAN, HELE
APPLICANT: KATTEVILD
APPLICANT: PALISSER,
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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STREET: **..
CITY: PALO ALTO
CTATE: CALIFORNIA
TISA
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APPLICANT:
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MEDIUM TYPE: Floppy disk
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RAE-VENTER PH.D., BARBARA
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AV-GAY, YOSSEF
NAN, HELENA M.
KATTEVILDER, ALFRED
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VENTION: AN OXIDO REDUCTASE ENZYME SYSTEM
VENTION: OBTAINABLE FROM P. CHRYSOGENUM, THE SET OF GENES ENCODING
VENTION: THE SAME AND THE USE OF OXIDO REDUCTASE ENZYME SYSTEMS OR
VENTION: GENES ENCODING THE SAME FOR INCREASING ANTIBIOTIC PRODUCTION
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US-09-562-702A-30
US-09-561-81BA-26
US-10-037-182-18
US-09-949-016-7247
US-09-949-016-7347
US-09-561-709B-1
US-09-561-81BA-12
US-09-561-81BA-12
US-09-561-81BA-12
US-09-949-016-5938
US-08-479-72B-2
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US-09-581-81BA-12
US-09-581-81BA-12
US-09-581-81BA-12
US-09-581-81BA-12
US-09-581-81BA-12
US-09-592-685-2
PCT-US95-02251-18
US-08-347-594A-2
US-08-347-594A-2
US-08-346-66053-6
PCT-US96-06053-6
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US-09-562-702A-32
US-09-561-818A-28
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US-10-188-495-67
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Sequence 67, Appl
Sequence 67, Appl
Sequence 28, Appl
Sequence 28, Appl
Sequence 30, Appl
Sequence 30, Appl
Sequence 18, Appl
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Sequence 12, Appli
Sequence 1, Appli
Sequence 10, Appli
Sequence 10, Appli
Sequence 5938, Appli
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Sequence 3, Appli
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US-08-274-043B-5
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-7622
TELEPAX: (415) 857-0663
TELEX: 380816 COOLEY PA
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                            COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" diskette
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/274,043B
FILING DATE: 12-JULY-1994
CLASSIFICATION + 435
PRIOR APPLICATION NUMBER: US 07/820,688
APPLICATION NUMBER: US 07/820,688
APPLICATION NUMBER: PCT/NL91/00101
FILING DATE: 18-JUNE-1991
APPLICATION NUMBER: PCT/NL91/00101
FILING DATE: 18-JUNE-1990
APPLICATION NUMBER: PCT/NL91/00101
FILING DATE: 18-JUNE-1990
APPLICATION NUMBER: PCT/NL91/00101
FILING DATE: 18-JUNE-1990
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APPLICANT: Aharono
APPLICANT: Van Der
APPLICANT: Cohen,
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Patent No.
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APPLICANT: Pallissa,
TITLE OF INVENTION: F
TITLE OF INVENTION: F
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TITLE OF INVENTION: C
TITLE OF INVENTION: C
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CITY: Palo Alto
STATE: CA
TATE: TY S.A.
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APPLICANT:
APPLICANT:
                                                  FILING DATE: 18-JUNE-1990 ATTORNEY/AGENT INFORMATION:
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ZIP: 943
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LOCATION:
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                 NAME: Rae-Venter, Barbara REGISTRATION NUMBER: 32,7
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 REFERENCE/DOCKET NUMBER:
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o. 5652132
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Av-Gay, Yossef
Nan, Helena M.
Kattevilder, Alfred
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pallissa, Harriet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bovenberg, Roelof
Schreiber, Rachel
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Cohen, Gerald
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                                                                                                                                                                                                                                                                                                                                                                                                                Law Offices of Barbara Rae-Venter O. Box 60039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 31; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     An oxido reductase enzyme system obtainable from P. chrysogenum, the set of genes encoding the same and the oxido reductase enzyme systems or genes encoding the same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    increasing antibiotic production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Roelof
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Best Local Similarity
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/274,043
FILING DATE: 12-JULY-1994
APPLICATION NUMBER: US 07/820,688
FILING DATE: 24-MARCH-1992
APPLICATION NUMBER: PCT/NL91/000101
FILING DATE: 18-JUNE-1991
APPLICATION NUMBER: EP 90201598.1
APPLICATION NUMBER: EP 90201598.1
FILING DATE: 18-JUNE-1990
ATTORNEY/AGENT INFORMATION:
NAME: Rae-Venter, Barbara
REGISTRATION NUMBER: 32,750
REFERENCE/DOCKET NUMBER: GBRO-024/02US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (415) 926-6200
TELEPAX: (415) 854-3713
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                   tent No.
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APPLICANT:
APPLICANT:
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TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Argaman, Anat
Av-Gay, Yossef
Nan, Helena M.
Kattevilder, Alfred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pallissa, Harriet
Pallissa, Harriet
VENTION: An oxido reductase enzyme system
VENTION: Obtainable from P. chrysogenum, from the use of
                                                                                                                                                                                                                                                                                                                                                               U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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d Hill Road, Ste.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoding the same and the use of oxido reductase enzyme systems or genes encoding the same for increasing antibiotic production
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Pred. No. 4.6e+05;
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Patent No. 5910435
GENERAL INFORMATION:
                                                        Matches
                                                                    Query Match
Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/687,276
FILING DATE: 25-JUL-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (415) 926-62
TELEFAX: (415) 854-3713
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                     TELEFAX: 608-251-9166
INFORMATION FOR SEQ ID NO:
                                                                                                                              TOPOLOGY: 1in MOLECULE TYPE: FRAGMENT TYPE:
                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27,386
REFERENCE/DOCKET NUMBER: 96
TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: protein
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                                                      Conservative
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internal
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                                                    100.0%; Score 31; DB 1;
100.0%; Pred. No. 4.6e+05;
tive 0; Mismatches 0;
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Pred. No. 4.6e+05,
0; Mismatches 0,
                                                                                    Length 4;
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                                                      Gaps
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RESULT 5
US-08-883-804-2
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                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21, Application US/08504538A Patent No. 6004746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100.0%; Score 31; DB 1; Length 4; Best Local Similarity 100.0%; Pred. No. 4.6e+05; Matches 4; Conservative 0; Mismatches 0; Indels
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APPLICANT: White, Carl W.
APPLICANT: Das, Kumuda C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: CONNELL, GARY J.
REGISTRATION NUMBER: 32,020
REFERENCE/DOCKET NUMBER: 2879
TELECOMMUNICATION INFORMATION:
TELEPHANE: 303/863-9700
TELEPHANE: 303/863-0223
                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/883,804
                                                           ZIP: 02110-2214
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                              NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: Clark & Elbing LLP
                                                                                                                                                                                                                                                                      APPLICANT: Brent, Roger
APPLICANT: McCoy, John M.
APPLICANT: Jessen, Timm H.
TITLE OF INVENTION: INTERACTION TRAP SYSTEMS FOR DETECTING
TITLE OF INVENTION: PROTEIN INTERACTIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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ADDRESSEE: Sheridan Ross P.C.
STREET: 1700 Lincoln Street, Suite 3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: USE OF THIOREDOXIN-LIKE MOLECULES FOR TITLE OF INVENTION: INDUCTION OF MASOD TO TREAT OXIDATIVE DAMAGE NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
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TOPOLOGY: linear
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                COUNTRY:
                                                                                                                                                                                            STREET:
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ER: 2879-40
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                                                                                                                                                                                                                                                                                                                                                                        US-09-579-883A-1
                                             GENERAL INFORMATION:
APPLICANT: BECKWITH, JONATHAN
APPLICANT: ASLUND, FREDRIK
APPLICANT: BESSETTE, PAUL H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1, Application US/09579883A Patent No. 6686443 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 14
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1
LENGTH: 4
TYPE: PRT
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Best Local
                                                                                                                   Sequence 2, Application US/09679705
Patent No. 6872563
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                                                                                                                                                                                                                                                                                                                                    Query Match
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              APPLICANT:
                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 31; DB 2; Best Local Similarity 100.0%; Pred. No. 4.6e+05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: RABENSTEIN, DALLAS
APPLICANT: SHI, TIESHENG
TITLE OF INVENTION: CHEMICAL REAGENTS FOR FORMATION OF DISULFIDE BONDS IN PEPTIDES
EILE REFERENCE: 407T-891100US
CURRENT APPLICATION NUMBER: US/09/579,883A
CURRENT FILING DATE: 2000-05-26
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FILING DATE: 07/20/95
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/278,082
FILING DATE: 07/20/94
ATTORNEY/AGENT INFORMATION:
NAME: Paul T. Clark
REGISTRATION NUMBER: 30,162
REGISTRATION NUMBER: 30,062
REGISTRATION NUMBER: 00/786/2:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 428-0200
TELEPAX: (617) 428-7045
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: synthetic peptide
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RITZ, DANIEL
LIM, JACKIE EUN-AH
                                GEORGIOU,
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                                GEORGE
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RESULT 9
US-08-883-804-4
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; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: variant motif
US-09-679-705-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/08883804
Patent No. 5985261
GRNERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: Das, Kumuda C.
TITLE OF INVENTION: USE OF THIORE
TITLE OF INVENTION: INDUCTION OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
  Query Match
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PRIOR FILING DATE: 1999-11-17
NUMBER OF SEQ ID NOS: 24
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PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: 60/163,939
PRIOR FILING DATE: 1999-11-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/679,705
CURRENT FILING DATE: 2000-10-05
                                                                                                                                                                                                                CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Connell, Gary J.
REGISTRATION NUMBER: 32,020
REFERENCE/DOCKET NUMBER: 2879
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR PRODUCTION OF DISULFIDE TITLE OF INVENTION: BOND CONTAINING PROTEINS IN HOST CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
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                                                                                                                                                           NFORMATION FOR SEQ ID NO:
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                                                                                                                                        SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
COMPUTER: LEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/883,804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                              MOLECULE TYPE:
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                                                                                  TOPOLOGY:
                                                                                                                     LENGTH:
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1700 Lincoln Street, Suite 3500
                                                                                                                     6 amino acids
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                                                           peptide
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  100.0%;
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  Score 31;
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DB 1;
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RESULT 11
US-08-627-907A-6
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Best Local Similarity
Thes 4; Conserve
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                                                                                                                                                                                                                                                                                                                                                 Sequence 6, Application US/08627907A
Patent No. 6060302
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APPLICATION: 514
ATTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNEY/ACTORNE
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Patent No.
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                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                        APPLICANT: HIRANO, NAOCO
APPLICANT: HIRAI, HISAMBATU
TITLE OF INVENTION: HUMAN PHOSPHOLIPASE C-ALPHA AND DNA
TITLE OF INVENTION: SEQUENCE ENCODING THE SAME
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NAME: CORDELL, Gary J.
REGISTRATION NUMBER: 32,020
REFERENCE/DOCKET NUMBER: 28
                                                                                                      NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
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ADDRESSEE: Sheridan Ross P.C.
STREET: 1700 Lincoln Street, Suite 3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: White, Carl W.
APPLICANT: Das, Kumuda C.
TITLE OF INVENTION: USE OF
TITLE OF INVENTION: INDUCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
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ADDRESSEE: c/o FISH & NEAVE
STREET: 1251 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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nilarity 100.0%;
Conservative 0
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Pred. No. 4.6e+05;
D; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Version
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                                                                                                ; OTHER INFORMATION: S. cerevisiae PDI thioredoxin active site
US-07-901-713A-3
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                                                                                                                                                                                                                                                                                                                                                              Sequence 3, Application US/07901713A

Patent No. 6291205

GENERAL INFORMATION:
APPLICANT: Tuite, Michael F.
APPLICANT: Freedman, Robert B.
APPLICANT: Markus, Henry Z.
APPLICANT: Schultz, Loren D.
APPLICANT: Montgomery, Donna L.
APPLICANT: Montgomery, Donna L.
APPLICANT: MILLS, Ronald W.
APPLICANT: BILLS, Ronald W.
TITLE OF INVENTION: METHOD FOR INCREASING PRODUCTION OF
TITLE OF INVENTION: DISULFIDE BONDED RECOMBINANT PROTEINS BY SACCAROMYCES
TITLE OF INVENTION: CEREVISIAE
FILE REFERENCE: 19460
                            Best
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Best Local S
                                                                                                                                                                                                                                    SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                  Query Match
                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/07/901,713A
CURRENT FILING DATE: 1992-06-12
NUMBER OF SEQ ID NOS: 23
SOPTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 18469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (212) 596-9090
INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING PATENT Release #1.0, Version #1.30
CURRENT APPLICATION DATA
APPLICATION NUMBER: US/08/627,907A
                                                                                                                                                                                                                   ENGTH: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: not relevant MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: SHGN-10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: JP 5-238402
FILING DATE: 24-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/JP94/01572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 22-SEP-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       y Match 100.0%; Score 31; DB 2; 1
Local Similarity 100.0%; Pred. No. 4.6e+05;
hes 4; Conservative 0; Mismatches 0;
y Match 100.0%; Score 31; DB 2; Length 6;
Local Similarity 100.0%; Pred. No. 4.6e+05;
hes 4; Conservative 0; Mismatches 0; Indels
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ZIP: 10020
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: HALEY Jr., Jam
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N
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       Gaps
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RESULT 13
US-08-571-985-27
Sequence 27, Application US/08571985
Patent No. 5783557
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                                                                                                                                                                                                                              RESULT 14
US-09-116-766-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/571,985
FILING DATE:
CLASSIFICATION: 514
PRIOR ADDITATION: 514
                                                                                                                                                                                                                                                                                                                                                                                   Best Loc
Matches
                                                                                                                                                                      Sequence 27, Applicat Patent No. 5968898 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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APPLICANT: Burstein, Yigal
APPLICANT: Trainin, Nathan
APPLICANT: Rycus, Avigail
TITLE OF INVENTION: THF-gamma2 Analogs and Pharmaceutical
TITLE OF INVENTION: Compositions Comprising Them
NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (810) 539-50:
INFORMATION FOR SEQ ID NO:
                                                                       APPLICANT: Burstein, Yigal
APPLICANT: Trainin, Nathan
APPLICANT: Rycus, Avigail
TITLE OF INVENTION: THF-gamma2 Analogs and Pharmaceutical
TITLE OF INVENTION: Compositions Comprising Them
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 01-JUL-1993
ATTORNEY/AGENT INFORMATION:
NAME: Kohn, Kenneth I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: IL 106214
                   NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kohn & As
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 30,955
REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 8 amino acids
TYPE: amino acid
TOPOLOGY: linear
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 STREET:
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Y: USA
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                                                                                                                                                                                                            Application US/09116766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Kohn & Associates
30500 No. 5783557thwestern Hwy., Suite 410
E: Kohn & Associates
30500 No. 5968898thwestern Hwy., Suite 410
                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 31; DI ilarity 100.0%; Pred. No. 4.6 Conservative 0; Mismatches
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COUNTRY:

USA

Farmington Hills

Michigan

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 34,
                                                                                                                                                                           ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Common OPERATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: (248) 539-5050
TELEPAX: (248) 539-5055
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
APPLICATION NUMBER: US/08/836,480
FILING DATE: 14-APR-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: SE 95\403256-8
FILING DATE: 14-OCT-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                    COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Bergstrand, Hakan
APPLICANT: Erickson, Tomas
APPLICANT: Karabelas, Kostas
APPLICANT: Lindvall, Magnus
APPLICANT: Sarinstrand, Bengt
TITLE OF INVENTION: NEW PEPTIDES WITH IMMUNOMODULATORY EFFECTS
NUMBER OF SEQUENCES: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 11 106214
FILING DATE: 01-JUL-1993
ATTORNEY/AGENT INFORMATION:
NAME: MONTGOMETY, Ilene N.
REGISTRATION NUMBER: 38,972
REFERENCE/DOCKET NUMBER: 2163.00050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                 CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 CGPC 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8 amino acids
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Pred. No. 4.6e+05;
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; MOLECULE TYPE: peptide
; SEQUENCE DESCRIPTION: SEQ ID NO: 34:
US-09-562-897-34
Query Match
Best Local Similarity
Thes 4; Conserva
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US-09-562-897-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; MOLECULE TYPE: peptide US-08-836-480-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 34, Application Patent No. 6228374
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 31; DB 2; Best Local Similarity 100.0%; Pred. No. 4.6e+05; Matches 4; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 617-542-8906 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                         TELEFAX: 617-542-8906 INFORMATION FOR SEQ ID NO: 34: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5970
TELEPAX: 617-542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FRASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/562,897
FILING DATE: 01-May-2000
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/836,480
FILING DATE: <Unknown>
PRIOR APPLICATION NUMBER: 08/836,480
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REGISTRATION NUMBER: 34,819
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER:
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                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sarnstrand, Bengt
TITLE OF INVENTION: NEW PEPTIDES WITH IMMUNOMODULATORY EFFECTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Bergstrand, Hakan
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    8 amino acids
amino acid
    XGY: linear

                                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                     TELEPHONE: 617-542-5070
                                                                                                                                                                                                          LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/09562897
                   100.0%; Score 31; DB 2; ilarity 100.0%; Pred. No. 4.6e+05; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Karabelas,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lindvall, Magnus
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                                                            Length 8;
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                       Indels
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                     0,
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US-08-750-142B-54
                                                                                                                                                                                          RESULT 18
US-09-160-513-217
                                                                                                                                                                                                                                                                         밁
                                                                                                                                                                                                                                                                                                         8
                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-750-142B-54
                                                                                                                                                 Sequence 217, Application US/09160513 Patent No. 6410775
                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 4; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 54, Application US/08750142B Patent No. 6228373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 617-542-8906
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/SE96/00:
FILING DATE: 22-MAR-1996
APPLICATION NUMBER: SE9501067-4
FILING DATE: 24-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/08/750,142B
FILING DATE: 25-NOV-1996
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bergstrand, Hakan
APPLICANT: Erikeson, Tomas
APPLICANT: Lindvall, Magnus
APPLICANT: Sarnstrand, Bengt
TITLE OF INVENTION: NEW PEPTIDES WITH
TITLE OF INVENTION: IMMUNOMODULATORY EFFECTS
                APPLICANT: Victoria, Edward J.
APPLICANT: Marquis, David M.
APPLICANT: Jones, David S.
APPLICANT: Yu, Lin
TITLE OF INVENTION: aPL IMMUNOREACTIVE PEPTIDES, CONJUGATES THEREOF AND METHODS NUMBER OF SEQUENCES: 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 55
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 617-542-5070
  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Boston
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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Pred. No. 73;
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RESULT 20
US-09-881-572A-25
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                                                                                                                                                       ; TYPE: PRT ; ORGANISM: Rhizobium leguminosarum US-09-230-041-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-230-041-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; MOLECULE TYPE: peptide US-09-160-513-217
                                                                                                                                                                                                               SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 31
LENGTH: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 6465179
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 31, Application US/09230041A Patent No. 6465179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                              Matches
                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                    APPLICANT: THIREOS, GEORGE
APPLICANT: KAPETZOPOULOS, DIMITRIS
TITLE OF INVENTION: DNA ENCODING AN ARTHROPOD CHITIN SYNTHASE
FILE REFERENCE: DNA ENCODING AN ARTHROPOD CHITIN SYNTH
CURRENT APPLICATION NUMBER: US/09/230,041A
CURRENT FILING DATE: 1999-01-19
EARLIER APPLICATION NUMBER: EP97108240.9
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ ID NOS: 42
COURTENER: ESEC ID NOS: 42
COURTENER: ESEC ID NOS: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: CATHERINE M. POLIZZI
REGISTRATION NUMBER: 40,130
REFERENCE/DOCKET NUMBER: 25231-20061.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/160,513
APPLICATION NUMBER: US/09/160,513
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STREET: 755 PAGE MILL ROAD
CITY: PALO ALTO
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          / Match
Local Similarity 100.0%; Pred. No.
nes 4; Conservative 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (650) 813-56
TELEFAX: (650) 494-0792
TELEX: 706141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 1
CLASSIFICATION:
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21 CGPC 24
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                                       2 CGPC 5
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                                                                           100.0%; Score 31; DB 2;
100.0%; Pred. No. 1.3e+02;
tive 0; Mismatches 0;
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; Sequence 26, Application US/10044359
Patent No. 6740743
; GENERAL INFORMATION:
APPLICANT: Herrman, Rafael
APPLICANT: Wong, James F.
APPLICANT: Lee, Jian-Ming
TITLE OF INVENTION: SCORPION TOXINS
FILE REFERENCE: BB1367 US NA
CURRENT FILING DATE: 2002-01-11
CURRENT FILING DATE: 2002-01-11
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                                                                                                                                                                                                                                             US-10-044-359-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-881-572A-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOPTWARE: PatentIn version 3.0
SEQ ID NO 5
LENGTH: 30
TYPE: PRT
ORGANISM: amphotropic murine leukemia virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Bupp, Keith;
APPLICANT: University of Medicine and Dentistry of New Jersey
TITLE OF INVENTION: Targeting Viral Vectors to Specific Cells
FILE REFERENCE: 601-1-095
CURRENT FILING DATE: 2002-03-05
PRIOR APPLICATION NUMBER: US/09/881,572A
CURRENT FILING DATE: 2002-03-05
PRIOR APPLICATION NUMBER: 60/212,239
PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.0
SEQ ID NO 25
LENGTH: 29
TYPE: DET
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Best Local Similarity
Matches 4; Conserv
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Patent No. 6762031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Bupp, Keith;
APPLICANT: University of Medicine and Dentistry of New Jersey
TITLE OF INVENTION: Targeting Viral Vectors to Specific Cells
FILE REFERENCE: 601-1-095
CURRENT APPLICATION NUMBER: US/09/881,572A
CURRENT FILING DATE: 2002-03-05
PRIOR APPLICATION NUMBER: 60/212,239
PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Roth, Monica;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Roth, Monica;
APPLICANT: Bupp, Keith;
PRIOR APPLICATION NUMBER: 09/599,416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 31; DB 2;
100.0%; Pred. No. 1.4e+02;
live 0; Mismatches 0;
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US-08-239-256-16
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SEQ ID NO 26
LENGTH: 35
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                                                            Matches
                                                                                          Query Match
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2000-06-22
PRIOR APPLICATION NUMBER: 60/140,227
PRIOR FILING DATE: 1999-06-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Androctonus mauretanicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                   NAME: MURASHIGE, KATE H.
REGISTION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-1500
                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: BOIME, IRVING
APPLICANT: MATZUK, MARTIN M.
APPLICANT: KEENE, JEFFREY L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: CTP EXTENDED FORM OF NUMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           y Match 100.0%;
Local Similarity 100.0%;
                                                                                                                                                                                                                                            TELEPHONE: (202) 887-0763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 2000 Peimis CITY: Washington, D.C.
                                                                             Local Similarity
                                                                                                                                           STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/239,256 FILING DATE: 06-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
                                                                                                                                                                         TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                          LENGTH:
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23
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                                                                                                                                                                                          38 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3: MORRISON & FOERSTER
2000 Pennsylvania Ave. N.W.
                                                            Conservative
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                                                                             100.0%; Score 31; DB 1; 100.0%; Pred. No. 1.6e+02;
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Pred. No. 1.6e+02;
                                                              Mismatches
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                                                              Gaps
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RESULT 24 US-09-244-583-20 ; Sequence 20, Application US/09244583

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APPLICANT: SOKER, SHAY

APPLICANT: MIAO, HUA-QUAN

APPLICANT: MIAO, HUA-QUAN

APPLICANT: MIAO, HUA-QUAN

TITLE OF INVENTION: NEURODILINS AND USE THEREOF IN METHODS FOR

TITLE OF INVENTION: DIAGNOSIS AND PROGNOSIS OF CANCER

FILE REFERENCE: 701039-48800

CURRENT APPLICATION NUMBER: US/09/583,638

CURRENT FILING DATE: 2000-05-30

PRIOR APPLICATION NUMBER: PCT/US98/26127

PRIOR APPLICATION NUMBER: 60/069,155

PRIOR APPLICATION NUMBER: 60/069,155

PRIOR APPLICATION NUMBER: 60/069,687

PRIOR FILING DATE: 1997-12-09

PRIOR PILING DATE: 1997-12-12

NUMBER: 00 SEQ ID NOS: 11

NUMBER: 05 SEQ ID NOS: 11
                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: PRT ; ORGANISM: Homo sapiens US-09-583-638-11
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US-09-244-583-20
                                                                                                                          US-09-579-420B-1
                                                                                                                                              RESULT 26
                   Sequence 1, Application US/09579420B Patent No. 6777534
GENERAL INFORMATION:
APPLICANT: Klagsbrun, Michael
APPLICANT: Soker, Shay
                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: P
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GENERAL INFORMATION:
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Sequence 11, App-
Sequence 11, App-
No. 6635421
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Best Local (
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Best Local Similarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/244,583
CURRENT FILING DATE: 1999-02-04
EARLIER APPLICATION NUMBER: 60/073,979
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 29
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 6479654
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APPLICANT: ANDREASON, GRAI
TITLE OF INVENTION: NOVEL FORMS OF THE ANGIOGENIC FACTOR
TITLE OF INVENTION: VASCULAR ENDOTHELIAL CELL GROWTH FAC
FILE REFERENCE: 240/086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: KLAGSBRUN, MICHAEL
APPLICANT: SOKER, SHAY
TITLE OF INVENTION: PEPTIDE ANTAGONISTS OF VASCULAR ENDOTHELIAL GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 4:
TYPE: PRT
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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CURRENT APPLICATION NUMBER: US/09/579,420B
CURRENT FILING DATE: 2000-05-25
PRIOR APPLICATION NUMBER: PCT/US98/26103
PRIOR FILING DATE: 1998-12-09
PRIOR REPLICATION NUMBER: 60/069,687
PRIOR FILING DATE: 1997-12-12
PRIOR APPLICATION NUMBER: 60/069,155
PRIOR APPLICATION NUMBER: 60/069,155
PRIOR APPLICATION NUMBER: 1997-12-09
NUMBER OF SEQ ID NOS: 19
SOPTWARE: PATENTIN VETSION 3.1
SEQ ID NO 1
                                                                                                                                                                                           US-09-513-999C-7252
; Sequence 7252, Application US/09513999C
; Patent No. 6783961
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US-09-579-420B-19
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-579-420B-19
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                                                                                                                                                                                                                                                                   RESULT 28
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 19
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins. Patent No. 6783961
FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
                                                                                                                                                                             GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity
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                                                                                                          APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: PETTIDE ANTAGONISTS OF VASCULAR ENDOTHELIAL GROWTH FACTOR FILE REFERENCE: 701039-04785-C CURRENT APPLICATION NUMBER: U09/579,420B CURRENT FILING DATE: 2000-05-25 PRIOR APPLICATION NUMBER: PCT/US98/26103 PRIOR FILING DATE: 1998-12-09 PRIOR FILING DATE: 1998-12-12 PRIOR FILING DATE: 1997-12-12 PRIOR FILING DATE: 1997-12-12 PRIOR APPLICATION NUMBER: 60/069,687 PRIOR FILING DATE: 1997-12-12 PRIOR FILING DATE: 1997-12-12 PRIOR FILING DATE: 1997-12-09
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APPLICANT: Soker, Shay
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
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100.0%; Pred. No. 2e+02;
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Pred. No.
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; FEATURE:
, NAME/KEY: UNSURE
; LOCATION: 27
; OTHER INFORMATION: Xaa=Arg or Ser or Thr
US-09-513-999C-5904
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US-09-513-999C-5905
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                                                                                                         Sequence 5905, Application US/09513999C Patent No. 6783961
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 5904
LENGTH: 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
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SOFTWARE: Patent.pm
SEQ ID NO 7252
                                                   TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins Patent No. 6783961
FILE REFERENCE: 59.US2.REG
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Best Local S
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24 PRIOR APPLICATION NUMBER: US 60/122,487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Homo sapiens
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Pred. No.
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US-09-630-273-2
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Sequence 2, Application US/09630273

Patent No. 6521416

GENERAL INFORMATION:
Birken et al, Steven
TITLE OF INVENTION: Determination Of The Amount Of hith Beta Core Fragment
TITLE OF INVENTION: In A Sample From A Subject And Uses Thereof
FILE REFERENCE: 54204-A-PCT-US
FILE REFERENCE: 54204-A-PCT-US
CURRENT APPLICATION NUMBER: US/09/630,273
CURRENT FILING DATE: 2000-08-01
PRIOR APPLICATION NUMBER: PCT/US99/02279
PRIOR FILING DATE: 1999-02-03
NUMBER OF SEQ ID NOS: 2
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15849)B
CURRENT APPLICATION NUMBER: US0/9/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 11828
LENGTH: 77
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Best Local Similarity
Matches 4; Conserv
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SOFTWARE: Patent.pm
SEQ ID NO 5905
LENGTH: 67
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Patent No. 6833447
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OTHER INFORMATION: >
FEATURE:
NAME/KEY: UNSURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 1999-02-26
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NAME/KEY: UNSURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Myxococcus xanthus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
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Local Similarity 100.0%; Pred. No. 2.7e+02;
hes 4; Conservative 0; Mismatches 0;
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Gregory J.
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Pred. No. 2.5e+02;
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RESULT 35
US-09-621-976-5629
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                                                                                                                                                                                                     ; TYPE: PRT
; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-697
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US-09-198-452A-697
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                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 697
LENGTH: 102
                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Griffais, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preve
TITLE OF INVENTION: and treatment of infection
FILE REFERENCE: 9710-003-999
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 697, Application US/09198452A Patent No. 6559294
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Best Local Similarity
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SOFTWARE: Patent.pm
SEQ ID NO 6168
LENGTH: 89
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTs and Encoded Human Proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
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1 Similarity 100.0%;
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                                                                 CGPC 31
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                                                                                                                                100.0%; Score 31; DB 2; 100.0%; Pred. No. 3.3e+02; cive 0; Mismatches 0;
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APPLICANT: Jobert, S. J.Y.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION. EST9 and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
CUURRENT APPLICATION NUMBER: US/09/621,976
CUURRENT FILING DAYE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
SOFTWARE: Patent.pm
SEQ ID NO 5629
LENGTH: 102
TYPE: PRT
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GENERAL INFORMATION:
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APPLICANT: Dumas Milne Edwards, J.B.
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                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201 487-8800
TELEPAX: 201 343-1684
                                                                                                                                                   TELEX: 133521
INFORMATION FOR SEQ ID NO:
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NAME/KEY: SIGNAL
LOCATION: -35..-1
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MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/241,853 FILING DATE: 12-MAY-1994 CLASSIFICATION: 435
                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 103 amino acids
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APPLICANT: Hanafusa, Hidesaburo
TITLE OF INVENTION: A TRANSMEMBRANE TYROSINE PHOSPHATASE
TITLE OF INVENTION: AND METHODS OF USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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STATE: New Jersey
                                                       STRANDEDNESS: single TOPOLOGY: linear
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100.0%; Pred. No. 3.3e+02;
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RESULT 38

US-09-583-110-3078

; Sequence 3078, Application US/09583110

; Patent No. 6699703

; GENERAL INFORMATION:
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Best Local Similarity
Matches 4; Conserv
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GENERAL INFORMATION:
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LENGTH: 103 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201 487-5800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Hanafusa, Hidesaburo
TITLE OF INVENTION: A TRANSMEMBRANE TYROSINE PHOSPHATASE
TITLE OF INVENTION: AND METHODS OF USE THEREOF
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
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APPLICATION NUMBER: US 08/241,853
FILING DATE: 12-MAY-1994
ATTORNEY/AGENT INFORMATION:
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TELEFAX: 133521
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STATE: New Jersey
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NO
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02-MAY-1997
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US-08-826-910-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Streptococcus pneumoniae US-09-583-110-3078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5210073-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 28-SEP-1990;SEQ ID NO:1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: YODOI, JUNJI; UCHIDA, ATSUSHI; TAGAYA, YUTAKA; MITSUI, AKIRA; HIRAKAWA, TADASHI
TITLE OF INVENTION: METHOD FOR TRREATING CANCER THERAPY; RADIATION DAMAGE OR ARTERIOSCLEROSIS USING HUMAN ADF
NUMBER OF SEQUENCES: 1
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: PATHO0-07A
CURRENT APPLICATION NUMBER: US/09/583,110
CURRENT FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/107,433
PRIOR FILING DATE: 1998-06-30
PRIOR APPLICATION NUMBER: US 60/085,131
PRIOR FILING DATE: 1998-05-12
PRIOR APPLICATION NUMBER: US 60/051,553
PRIOR APPLICATION NUMBER: US 60/051,553
PRIOR FILING DATE: 1997-07-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 5210073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 3078
LENGTH: 104
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                    Sequence 3, Application US/08826910 Patent No. 5919657
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Best Local Similarity
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                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
APPLICANT: Shah, Purvi
TITLE OF INVENTION: THIOREDOXIN-LIKE PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Lynn Doucette-Stamm et al.
TITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Relating
TITLE OF INVENTION: Pneumoniae for Diagnostics and Therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
                                                                                                COUNTRY: UZIP: 94304
                                                                                                                                                      STREET: 3174 POI
CITY: Palo Alto
                                                                                                                                                                          ADDRESSEE: Incyte Pharmaceuticals, Inc STREET: 3174 Porter Drive
                                                                                                                                         STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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100.0%; Pred. No. 3.4e+02;
tive 0; Mismatches 0;
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CURRENT APPLICATION DATA:

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US-08-826-910-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
APPLICANT: Shah, Purvi
TITLE OF INVENTION: THOREDOXIN-LIKE PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                              ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
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IMMEDIATE SOURCE:
LIBRARY: GenBar
CLONE: 339649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                   APPLICATION NUMBER: US/08/826,910
FILING DATE: Herewith
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 105 amino acids
                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
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STRANDEDNESS: sir
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
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                                                TELEFAX:
                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                        COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94304
                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          32 CGPC 35
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                                                  415-845-4166
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                                                                                                                                                                                                                                                                                                                    IBM Compatible
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                                                                                                   PF-0268 US
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Pred. No. 3.4e+02;
; Mismatches 0;
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LIBRARY: GenBa
CLONE: 453972
US-08-826-910-4
; LENGTH: 105
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-513-999C-8037
                                                                                                                                                                                                                                                                                                                                                                           US-09-513-999C-8037
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US-09-601-144-68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 68
LENGTH: 105
TYPE: PRT
                                                                                                                                                                                                                         APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
                                                                                                                                                                                                                                                                                                                  Sequence 8037, Application US/09513999C Patent No. 6783961 GENERAL INFORMATION:
                                                                           SEQ ID NO 8037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Patent No. 6566514
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Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
                                                                                      FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 74
SOFTWARE: Patentin Ver. :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: OLIGONUCLEOTIDE SEQUENCES COMPLEMENTARY TO THIOREDOXIN TITLE OF INVENTION: AND THIOREDOXIN REDUCTASE GENES AND METHODS OF USING TITLE OF INVENTION: SAME TO MODULATE CELL GROWTH FILE REFERENCE: 683-112US-A CURRENT APPLICATION NUMBER: US/09/601,144

CURRENT FILING DATE: 2000-10-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1998-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Wright, Jim A. APPLICANT: Young, Aiping H. APPLICANT: Lee, Yoon S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
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100.0%; Pred. No.
tive 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 31; DB 1; 100.0%; Pred. No. 3.4e+02;
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Best Local Similarity
Marches 4; Conserva
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US-08-386-729A-10
                                                             US-08-386-729A-10
                                                                                                                                FILING DATE: 18-JUNE-1990
ATTORNEY/AGENT INFORMATION:
NAME: Rae-Venter, Barbara
REGISTRATION NUMBER: 32,750
REFERENCE/DOCKET NUMBER: GBRO-024/02US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 926-6200
TELEPHONE: (415) 936-4210
TELEPAX: (415) 854-3713
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 106 amino acids
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 10,
                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/274,043
FILING DATE: 12-JULY-1994
APPLICATION NUMBER: US 07/820,688
FILING DATE: 24-MARCH-1992
APPLICATION NUMBER: PCT/NL91/000101
FILING DATE: 18-JUNE-1991
PILING DATE: 18-JUNE-1991
APPLICATION NUMBER: EP 90201598.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Aharon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: An oxido reductase enzyme system
TITLE OF INVENTION: obtainable from P. chrysogenum, the set of genes
TITLE OF INVENTION: encoding the same and the use of oxido reductase enzyme
TITLE OF INVENTION: systems or genes encoding the same for increasing antibiotic
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSE: Wail
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Argaman,
APPLICANT: Av-Gay, y
APPLICANT: Nan, Hele
APPLICANT: Kattevild
APPLICANT: Pallissa,
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" diskette
COMPUTER: IBM PC compatible
COMPUTER: PATEN: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/386,729A
FILING DATE: 10-FEB-1995
CLASSIFICATION: 435
CLASSIFICATION: A35
CLASSIFICATION: A35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
                                                                           MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 2882 Sand
CITY: Menlo Park
STATE: CA
COUNTRY: U.S.A.
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                                                                                                LENGTH: 106 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               32 CGPC 35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Weil, Gotshal & Manges
2882 Sand Hill Road, Ste. 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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Bovenberg, Roelof A. L. Schreiber, Rachel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Argaman, Anat
Av-Gay, Yossef
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100.0%;
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Score 31; DB 1;
Pred. No. 3.4e+02;
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                  Length 106;
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APPLICANT: Abernethy, Nevin
APPLICANT: Conrust, Rene
APPLICANT: Conrust, Rene
APPLICANT: Murison, Greg
TITLE OF INVENTION: Compounds isolated from stromal cells
TITLE OF INVENTION: and methods for their use
FILE REFERENCE: 1100.1037c1
CURRENT APPLICATION NUMBER: US/09/383,586
CURRENT FILING DATE: 1999-08-26
NUMBER OF SEQ ID NOS: 38
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 13
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                                                                                                                               ; TYPE: PRT
; ORGANISM: Mouse
US-09-823-038A-13
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US-09-823-038A-13
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Best Local Similarity
Matches 4; Conserva
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US-09-383-586-13
                                                                                                                                                                                            APPLICANT: Kumble, Anand
APPLICANT: Murison, Greg
TITLE OF INVENTION: Compositions Isolated From Stromal Cells
TITLE OF INVENTION: and Methods For Their Use
TITLE REFERENCE: 11000.1037c3
CURRENT APPLICATION NUMBER: US/09/823,038A
CURRENT FILING DATE: 2001-07-09
NUMBER OF SEQ ID NOS: 61
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 13, Application US/09823038A Patent No. 6797271
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                                                                Matches
                                                                                Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Strachan, Lorna APPLICANT: Sleeman, Matthew
                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
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TYPE: PRT
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                                 2 CGPC 5
                                                              2h 100.0%; Score 31; DB 2; Similarity 100.0%; Pred. No. 3.4e+02
4; Conservative 0; Mismatches 0
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100.0%; Pred. No. 3.4e+02;
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RESULT 47
US-07-745-382-22
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US-07-921-848-22
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                                                                                                                                                                                                                                                              Patent No. 5292646
GENERAL INFORMATION:
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Best Local (
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                                                                                                                                                                                                                                                                                                Sequence 22,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
TYPE: AMINO ACID
TYPE: AMINO ACID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: CBerr, Luann
REGISTRATION NUMBER: 31,822
REGISTRATION NUMBER: GI5188A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Peptide and Protein Fusions To
TITLE OF INVENTION: Thioredoxin and Thioredoxin-Like Molecules
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc.
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: McCoy, John
APPLICANT: LaVallie, Edward
                                                                                                                                                          NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                      APPLICANT: McCoy, John
APPLICANT: LaVallie, Edward
TITLE OF INVENTION: Peptide and Protein Fusions To
TITLE OF INVENTION: Thioredoxin and Thioredoxin-Like Molecules
                                                                                         STREET: 0, CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: Massachusetts COUNTRY: U.S.A.
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                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
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                                                                                                                                           ADDRESSEE:
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5292646
                                                                                         Massachusetts
                                                                                                                       E: Genetics Institute,
87 CambridgePark Drive
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                                                                         U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 31; DB 1; ilarity 100.0%; Pred. No. 3.5e+02; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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Best Local Similarity
Matches 4; Conserv
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APPLICATION NUMBER: US 07/652,531
FILING DATE: 06-FEB-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/745,382
FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: CBGIT, LUMIN
REGISTRATION NUMBER: 31,822
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION UNMER: 33,544
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEFAX: (617) 876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (617) 876-58: INFORMATION FOR SEQ ID NO:
TELEFAX: (617) 876-5851 INFORMATION FOR SEQ ID NO:
                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                         STREET: b, STREET: Cambridge CITY: Cambridge CTATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith, Elizabeth
APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 109 amino acids
TYPE: AMINO ACID
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CLASSIFICATION:
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SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                  SOFTWARE:
                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A. ZIP: 02140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Grant, Kathleen
LaVallie, Edward R.
                                                                                                                                                                                                                                PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                              Genetics Institute,
                                                                                                                                                                                                                                                                                           Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PEPTIDE AND PROTEIN FUSIONS TO THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
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RESULT 51
US-08-165-301A-28
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                                                                                                                                                                                                                                          US-08-165-301A-26
Sequence 28, Application US/08165301A Patent No. 5646016
                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                      Matches
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Best Local (
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                                                                                                                                                                                                                                                                                   TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
TYPE: amino acid
TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CHAPTION NUMBER: US/08/165,301A
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 87 Cambridge
CITY: Cambridge
STATE: Massachusetts
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                                                                                                     33 CGPC 36
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                                                                                                                                                                    100.0%; Score 31; I ilarity 100.0%; Pred. No. 3. Conservative 0; Mismatches
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US-08-810-436-22
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                                                                                                                                                                                                                                                                                                                                                     Patent No. 6143524
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                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
OPERATING SYSTEM: PC-DOS
                                                                                                                                                                                                           APPLICANT: Grant, Kathleen

APPLICANT: LaVallie, Edward R.

APPLICANT: LaVallie, Edward R.

TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO

TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN LIKE MOLECULES, AND MODIFIED

TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/165,301A
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TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                         NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                   APPLICANT: McCoy, John APPLICANT: DiBlasio-Sm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: McCoy,
APPLICANT: DiBlasi
APPLICANT: Grant,
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                                                                                    COUNTRY:
                                                                                                                         STREET: 87 Cambı
CITY: Cambridge
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
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                                                                                                         Massachusetts
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                                                                                                                                      RE: Genetics Institute,
87 CambridgePark Drive
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87 CambridgePark Drive
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                                                                                      U.S.A.
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Grant, Kathleen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 31; DB 1; 100.0%; Pred. No. 3.5e+02;
 PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                       Elizabeth
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CLASSIFICATION:

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/165,301
APPLICATION NUMBER: 08/165,301
PILING DATE: 10-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI 518
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-5851
TELEPAX: 700 ID NO: 26:
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US-08-810-436-26
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                                                                                                                                                                                                    CLODPY disk
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, VE
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/810,436
FILING DATE: 04-MAR-1997
CLASSIFICATION: 435
PRIOR APPLICATION: 435
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith, Elizabeth
APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES,
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/165,301 FILING DATE: 10-DEC-1993 ATTORNEY, AGENT INFORMATION:
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 87 Cambridge
CITY: Cambridge
STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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6143524
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04-MAR-1997
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                                                                                              GI 5188D
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; LENGTH: 109 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-810-436-26
片
                                                                                                                            TOPOLOGY: linear; MOLECULE TYPE: protein US-08-810-436-28
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Best Local Similarity 100
Matches 4; Conservative
                                                               Matches
                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/810,43
FILING DATE: 04-MAR-1997
CLASSIFICATION: 435
FRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/165,301
FILING DATE: 10-DEC-1993
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                              TELEFAX: (617) 876-58: INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                           NAME: Meinert, M. C.
REGISTAN NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELERHONE: (617) 876-1170
                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PATENTIA RELEASE #1.0, Version #1.25
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESSE: Genetics Institute,
ADDRESSEE: 87 CambridgePark Drive
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TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
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APPLICANT: LaVallie, Edward R.
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CITY: Cambridge
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33 CGPC 36
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                                                             4; Conservative
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04-MAR-1997
V. 17-
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                                                                              100.0%; Score 31; DB 2;
100.0%; Pred. No. 3.5e+02;
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BER: GI
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RESULT 55 US-09-485-885-17

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US-09-248-796A-19911
; Sequence 19911, Application US/09248796A
; Patent No. 6747137
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                                                                  RESULT 57
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SEQ ID NO 17
LENGTH: 109
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Best Local Similarity 100
Matches 4; Conservative
GENERAL INFORMATION:
                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 4; Conserv
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OTHER INFORMATION: amino acid of thioredoxin
-09-166-966E-10
                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 10
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APPLICANT:
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Patent No. 634222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: HSEU, TZONG-HSIUNG
TITLE OF INVENTION: CHIMERIC PROTEINS WITH A CELLULOSE BINDING DOMAIN
FILE REFERENCE: 32250-159660
CURRENT APPLICATION NUMBER: US/09/166,966E
CURRENT FILING DATE: 1998-10-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 198-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: CHEN, DAVID CHANHAN APPLICANT: HU, NIEN-TAI
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TITLE OF INVENTION: Vaccine
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                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: TW 86114750 PRIOR FILING DATE: 1997-10-08
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                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 10
TYPE: PRT
                                                                                                                                                                                                                                                                                                                           FEATURE:
NAME/KEY: thioredoxin
                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Unknown
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                                                                                                                          33 CGPC 36
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HSEU, TZONG-HSIUNG
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Delisse, Anne-Marie Eva Fernande
Gerard, Catherine Marie Ghislaine
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100.0%; Pred. No. 3.5e+02;
cive 0; Mismatches 0;
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RESULT 58
PCT-US94-14179-22
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Best Local Similarity 100.0%;
Matches 4; Conservative 0
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SEQ ID NO 19911
LENGTH: 109
TYPE: PRT
    Matches
                                       Query Match
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAL
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR PELICATION NUMBER: US 60/096,409
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
PRIOR FILING DATE: 1998-08-13
                                                                                                                                                                                                 TELEFAX: (617) 876-58
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve
CURRENT APPLICATION NUMBER: PCT/US94/14179
APPLICATION NUMBER: PCT/US94/14179
                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                               TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 876-1170
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y Match 100.0%; Score 31; DB 4; Cocal Similarity 100.0%; Pred. No. 3.5e+02; hes 4; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
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                                                                                                                                                            109 amino acids
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                                                                                                                                                                                             7) 6/-
) 876-5851
NO: 22:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Edward R.
                                                                                                                                                                                                                                                                                               33,544
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Pred. No. 3.5e+02;
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                                         DB 4; Length 109;
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RESULT 60
PCT-US94-14179-28
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; Sequence 26, Application PC/TUS9414179
; GENERAL INFORMATION:
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                                                                                                                                                                  Sequence 28, Application PC/TUS9414179
GENERAL INFORMATION:
APPLICANT: McCOY, John
APPLICANT: DiBlasio-Smith, Elizabeth
APPLICANT: DiBlasio-Smith, Elizabeth
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, ITTLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES, ITTLE OF INVENTION
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Best Local Similarity
Matches 4; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATECHTIN Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/14179
FILING DATE:
CLACETOTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: GI 5188D
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-5851
NFORMATION FOR SEQ ID NO: 26:
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NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
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APPLICANT: LAVAILE, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
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CORRESPONDENCE ADDRESS:
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                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                          NUMBER OF SEQUENCES:
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CITY: Cambridge
STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
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                                               STREET:
                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               33 CGPC 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
Cambridge
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                                   B: Genetics Institute, Inc
87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                     AND MODIFIED
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US-09-621-976-4028
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NAME: Meinert, M. C.

REGISTRATION NUMBER: 33,544

REFERENCE/DOCKET NUMBER: GI 5

TELECOMMUNICATION INFORMATION:

TELEPHONE: (617) 876-1170

TELEPAX: (617) 876-5851

INFORMATION FOR SEQ ID NO: 28:

SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 19335
SOFTWARE: Patent.pm
SEQ ID NO 4028
                                                              Matches
                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 4028, App
Patent No. 6639063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Dumas Milne Edwards, APPLICANT: Jobert, S. APPLICANT: Giordano, J.Y.
                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: ESTs and Encoded Human Proteins. FILE REFERENCE: GENSET.054PR2
                                                                                                                                          ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: SIGNAL
LOCATION: -102..-1
                                                                                                                                                                                                            LENGTH: 110
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: PCT/US94/14179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: Massachusetts
74 CGPC
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                                2 CGPC 5
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                                                              4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09621976
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 77
                                                                             100.0%;
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100.0%; Pred. No. 3.5e+02;
tive 0; Mismatches 0;
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                                                                               Score 31; DB 2;
Pred. No. 3.5e+02;
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                                                                Mismatches
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                                                                                            Length 110;
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RESULT 62 US-09-264-419C-6 ; Sequence 6, Application US/09264419C ; Patent No. 6174682 ; GENERAL INFORMATION:

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; OTHER INFORMATION: Description of Artificial Sequence: consensus; OTHER INFORMATION: sequence US-09-264-419C-6
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                                                                                                                                                                                                                                                      US-09-107-532A-6739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; ORGANISM: Staphylococcus epidermidis US-09-134-001C-3859
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 3859
LENGTH: 111
                                                                                                                                                                                              Sequence 6739, Application US/09107532A
Patent No. 6583275
GENERAL INFORMATION:
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Best Local Similarity
Matches 4; Conser
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: WNI-076
CURRENT APPLICATION NUMBER: US/09/264,419C
CURRENT FILING DATE: 1999-03-08
NUMBER OF SEQ ID NOS: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/134,001C CURRENT FILING DATE: 1998-08-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: LYNIN DOUGETTE-Stamm et al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: GTC-007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Khodadoust, Mehran
TITLE OF INVENTION: No. 6174682el Thioredoxin Family Active Site Molecules and
TITLE OF INVENTION: Therefor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1997-08-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/064,964
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1997-11-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS
                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                           APPLICANT: Lynn A Doucette-Stamm TITLE OF INVENTION: NUCLEIC ACID ENTEROCOCCUS
                                                                                                                                                                                                                                                                                                                                36 CGPC 39
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STATE: Massachusetts
COUNTRY: USA
ZIP: 02354
                                                     CITY: Waltham
                                                                      STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                  100 Beaver Street
                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 31; DB 2; 100.0%; Pred. No. 3.5e+02;
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Pred. No. 3.5e+02;
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                                                                                                                                         and David Bush
AND AMINO ACID SEQUENCES RELATING TO
FARCIUM FOR DIAGNOSTICS AND THERAPEUT
                                                                                         CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 111;
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GENERAL INFORMATION:
APPLICANT: HELLX RESEARCH INSTITUTE
FILE OF INVENTION: No. 6943241el full length cD:
FILE REFERENCE: H1-A0105
CURRENT APPLICATION NUMBER: US/10/104,047
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER:
PRIOR FILING DATE:
NUMBER OF SEQ ID NOS: 4096
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 3199
LENGTH: 113
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US-10-104-047-3199
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                                                                                                                                                     ; ORGANISM: Homo sapiens
US-10-104-047-3199
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                                                                         Matches
                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3199, Application US/10104047 Patent No. 6943241
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Best Local Similarity
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                                                                                                                                                                                           TYPE: PRT
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APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
APPLICATION NUMBER: 60/051571
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/107,532A
FILING DATE: 30-Jun-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc feature
LOCATION: (B) LOCATION 1...112
SEQUENCE DESCRIPTION: SEQ ID NO: 6739:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: protein HYPOTHETICAL: YES ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
TELEPHONE: (781)893-5007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
COMPUTER: PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          36 CGPC 39
                                     2 CGPC 5
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                                                                         4; Conservative
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SOFTWARE: ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (781)893-8277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Ariniello, Pamela Deneke REGISTRATION NUMBER: 40,489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Enterococcus faecium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                         100.0%; Score 31; DB 2;
100.0%; Pred. No. 3.6e+02;
tive 0; Mismatches 0;
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Pred. No. 3.5e+02;
); Mismatches 0;
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                                                                                                               Length 113;
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                                                                         0;
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US-08-918-288-71

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RESULT 67
US-09-282-357-71
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No. 6238890
                                                                                                                                                                          Patent No. 6242580
GENERAL INFORMATION:
                                                                                                                                                                                               Sequence 71, Application US/09282357 Patent No. 6242580
                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
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APPLICANT: BOIME, Irving
APPLICANT: MOYLE, William R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 08/853,524
APPLICATION NUMBER: 08/853,524
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: 08/199,382
FILING DATE: 18-FEB-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: FASTEM: DOS
SOFTWARE: FASTESQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatik
OPERATING SYSTEM: DOS
                                              NUMBER OF SEQUENCES: 83
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 114 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: 202-887-1500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: SINGLE-CHAIN FORMS OF THE TITLE OF INVENTION: GLYCOPROTEIN HORMONE QUARTET
                                                                                                     TITLE OF INVENTION: SINGLE-CHAIN FORMS OF THE TITLE OF INVENTION: GLYCOPROTEIN HORMONE QUARTET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: ZUUC
CITY: Washington
                                                                                                                                         APPLICANT: BOIME, Irving APPLICANT: MOYLE, William R.
STREET: ZUUC
STREET: ZUUC
STREET: ZUUC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
TOPOLOGY: lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Murashige, Kate H
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                          90 CGPC 93
                                                                                                                                                                                                                                                                                                                                            2 CGPC 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
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                              2000 Pennsylvania Avenue, NW, suite 5500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         114 amino acids
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                                                                                                                                                                                                                                                                                                                                                                               Conservative
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100.0%; P
tive 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09/282,357
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                                                                                                                                                                                                                                                                                                                                                                            Score 31; DB 2; 1
Pred. No. 3.6e+02;
); Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                               Length 114;
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US-09-897-425-41
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; ORGANISM: Arabidopsis thaliana US-09-897-425-41
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                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/897,425
CURRENT FILING DATE: 2001-07-03
PRIOR APPLICATION NUMBER: 09/210,843
PRIOR FILING DATE: 1998-12-18
PRIOR APPLICATION NUMBER: 08/846,021
PRIOR PILING DATE: 1997-04-25
PRIOR APPLICATION NUMBER: 08/366,783
PRIOR APPLICATION NUMBER: 08/366,783
PRIOR PILING DATE: 1994-12-30
PRIOR APPLICATION NUMBER: 08/142,418
PRIOR PILING DATE: 1993-11-16
PRIOR APPLICATION NUMBER: 08/1659,835
PRIOR APPLICATION NUMBER: 07/659,835
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100.0%; Score 31; DB 2;
Best Local Similarity 100.0%; Pred. No. 3.6e+02;
Matches 4; Conservative 0; Mismatches 0;
                                                                                  SOFTWARE: Pa
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Patent No. 675004
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: MOLONEY, MAURICE M.
APPLICANT: DALMIA, BIFIN K.
TITLE OF INVENTION: PREPARATION OF THIOREDOXIN AND THIOREDOXIN REDUCTASE
TITLE OF INVENTION: PROTEINS ON OIL BODIES
FILE REFERENCE: 034547/0106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX:
INFORMATION FOR SEQ ID NO:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/918
FILING DATE: 25 AUG-1997
APPLICATION NUMBER: 08/85:
FILING DATE: 09-MAY-1997
                                                                                                                                NUMBER OF SEQ ID NOS: 55
                                                                                                                                                     PRIOR FILING DATE: 1991-02-22
                                       TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Murashige Kate H
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-887-1500
                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 114 amino acid
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MEDIUM TYPE: Diskette
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SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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APPLICATION NUMBER: 08/199,382
FILING DATE: 18-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
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                                                                  114
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                                                                                                         PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/09897425
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RESULT 70
US-09-786-715-11
; Sequence 11, Application US/09786715
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                                                                                                                                   Query Match
Best Local Similarity luv.
"""hes 4; Conservative
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PATENT NO. 6800744
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID
SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE
THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local
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INFORMATION FOR SEQ ID NO: 2930:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/085131
FILING DATE: May 12, 1998
APPLICATION NUMBER: 60/05153
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-011
TELECHONE: (781)893-5007
TELECHONE: (781)893-8077
TELECAX: (781)893-8277
                                                                                                                                                                                                                                                   NAME/KEY: misc_feature
LOCATION: (B) LOCATION 1...114
SEQUENCE DESCRIPTION: SEQ ID NO: 2930:
                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
HYPOTHETICAL: YES
ORIGINAL SOURCE:
ORGANISM: Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/107,433 FILING DATE: 30-Jun-1998 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
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MEDIUM TYPE: CD/ROM ISO9660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME
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                                                                                         38 CGPC 41
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                                                                                                                    2 CGPC 5
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                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 114 amino acids TYPE: amino acid
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OPERATING SYSTEM: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100 Beaver Street
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                                                                                                                                                                            100.0%; Score 31; DB 2; 100.0%; Pred. No. 3.6e+02;
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Pred. No. 3.6e+02;
); Mismatches 0;
                                                                                                                                                               Mismatches
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                                                                                                                                    ; SOFTWARE: Microsoft Office 97
; SEQ ID NO 11
; LENGTH: 114
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
US-09-786-715-11
                                                                                                                                                                                                                                      APPLICANT: Allen, Steve
APPLICANT: Lu, Albert
APPLICANT: Lu, Albert
APPLICANT: Thorpe, Cathy
TITLE OF INVENTION: Thioredoxin H Homologs
FILE REFERENCE: BB-1246
CURRENT APPLICATION NUMBER: US/09/786,715
CURRENT FILING DATE: 2001-06-25
PRIOR FILING DATE: September 8, 1998
NUMBER OF SEQ ID NOS: 13
                                                                                  Query Match 100.0%;
Best Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. GENERAL IN
                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION:
40
                                2 CGPC 5
                                                                  4; Conservative
CGPC 43
                                                                   <u>.</u>
                                                                                   Score 31; DB 2; Pred. No. 3.6e+02;
                                                                   Mismatches
                                                                                                  Length 114;
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Gaps

0,

Search completed: February 23, 2006, 00:43:45 Job time: 44.5 secs

Cologn Madrid (USOTO)

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Result
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scoring table:
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1:
      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score
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      protein search, using
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length: 2000000000
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Match
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Maximum |
Listing
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      0000000000000000000000
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/cgn2 6/ptcodata/1/pubpaa/US07_pubcoMB.pep:*
/cgn2 6/ptcodata/1/pubpaa/US08_PUBCOMB.pep:*
/cgn2 6/ptcodata/1/pubpaa/US09_PUBCOMB.pep:*
/cgn2-6/ptcodata/1/pubpaa/US10A_PUBCOMB.pep:*
/cgn2-6/ptcodata/1/pubpaa/US10A_PUBCOMB.pep:*
/cgn2-6/ptcodata/1/pubpaa/US111_PUBCOMB.pep:*
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Match 100%
first 1000 summaries
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      GenCore version (c) 1993 - 2006
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US-10-286-516A-11
US-10-34-319A-1
US-10-660-118A-1
US-10-660-118A-1
US-10-628-391-2
US-10-776-933-14
US-10-304-287-4
US-10-290-072-233
US-10-660-118A-3
US-10-660-118A-3
US-10-660-118A-3
US-10-660-118A-3
US-10-660-118A-3
US-10-461-224-1
US-10-461-224-1
US-10-461-224-1
US-10-484-251-11
US-09-572-404B-970
US-10-044-844-217
US-10-044-844-217
US-10-044-844-217
US-10-053-339-10
US-10-053-903-98
US-10-053-903-98
US-10-053-903-98
US-10-053-903-98
US-09-881-572A-25
US-09-881-572A-5
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Sequence 8, Appli Sequence 7, Appli Sequence 155, App Sequence 9, Appli Sequence 20, Appli Sequence 5, Appli	quence quence quence	quence	quence	quence	quence	quence	Sequence 34096, A	Sequence 39815, A	Sequence 295, App Sequence 334048,	Sequence 64, Appl	Sequence 3343, Ap Sequence 106, App	Sequence 389, App	Sequence 182799,	Sequence 53092, A Sequence 2216, Ap	Sequence 34, Appl Sequence 36, Appl	Sequence 72, Appl Sequence 40, Appl	Sequence 52, Appl Sequence 60, Appl	Sequence 50, Appl	Sequence 70, Appl Sequence 38, Appl Seguence 55347	Sequence 63276, A	Sequence 4881, Ap	Sequence 66365, A	Sequence 87, Appl Sequence 8, Appli	Sequence 41147, A Sequence 7, Appli	Sequence 105, Appl Sequence 56, Appl	Sequence 4, Appli	10	5 1 1 1 1 1 1	6,	ω,ω	1,	58, App	1 6	5826, A	5 6	
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US-10-286-516A-11
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                Sequence 1, Application US/10234319A
Publication No. US20030109457A1
GENERAL INFORMATION:
APPLICANT: Atlas, Daphne
TITLE OF INVENTION: MOLTI-COMPONENT ANTIOXIDANT COMPOUNDS, PHARMACEUTICAL COMPOSITION
TITLE OF INVENTION: CONTAINING SAME AND THEIR USE FOR REDUCING OR PREVENTING OXIDATI
PILE REFERENCE: 24890
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                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
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APPLICANT: Walker, David H.
TITLE OF INVENTION: Ehrlichia Disulfide Bond Formation
TITLE OF INVENTION: Proteins and Uses Thereof
FILE REFERENCE: D6428
CURRENT APPLICATION NUMBER: US/10/286,516A
CURRENT FILING DATE: 2002-11-01
PRIOR APPLICATION NUMBER: US 60/335,611
PRIOR FILING DATE: 2001-11-01
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FEATURE:
OTHER INFORMATION: Escherichia coli conserved cysteine
OTHER INFORMATION: motif in active site of thioredoxin
OTHER INFORMATION: protein
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US-10-343-650A-448
US-10-1343-650A-448
US-10-819-316-441
US-10-099-322-58
US-10-369-493-21120
US-10-369-493-21120
US-10-1012-896-1011
US-10-144-678A-1011
US-10-369-493-21509
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US-10-435-696-52
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US-09-878-178-2235

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US-10-066-543-1422
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US-10-902-531-18
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Sequence 14, Appl
Sequence 326082,
Sequence 162, App
Sequence 2235, Ap
Sequence 2235, Ap
Sequence 2235, Ap
Sequence 2235, Ap
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Sequence 441, Appl
Sequence 441, Appl
Sequence 448, Appl
Sequence 58, Appl
Sequence 58, Appl
Sequence 58, Appl
Sequence 51120, Appl
Sequence 1011, Appl
Sequence 1011, Appl
Sequence 1011, Appl
Sequence 23509, Appl
Sequence 23509, Appl
Sequence 54715, Appl
Sequence 5
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US-10-369-094-1
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; LOCATION: (4)...(4)
; OTHER INFORMATION:
US-10-234-319A-1
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US-10-660-118A-1
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
LENGTH: 4
                                                                                                                                                                             Sequence 1, Application US/10660118A
Publication No. US20040131606A1
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TYPE: PRT
ORGANISM: homo
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ORGANISM: Artificial sequence
FEATURE:
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LOCATION: (1)...(1)
OTHER INFORMATION: ACETYLATION
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GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum FILE REFERENCE: 2879-98
CURRENT APPLICATION NUMBER: US/10/660,118A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Richon, Victoria M.
APPLICANT: Marks, Paul A.
APPLICANT: Rifkind, Richard A.
APPLICANT: Butler, Lisa M.
TITLE OF INVENTION: Method of Treating TRX Mediated
FILE REFERENCE: 3254-1005-001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/369,094
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: U.S. 60/357,383
PRIOR FILING DATE: 2002-02-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 31; DB 4; 100.0%; Pred. No. 1.7e+06;
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100.0%; Pred. No. 1.7e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn version 3.2 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/10628391 Publication No. US20040146526A1 GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
                                                                                                                                                                                                                                                                                 Sequence 149, Application US/10776933
Publication No. US20040241717A1
                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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Best Local
                                                                   APPLICANT: HANSEN, BO
APPLICANT: THRUE, CHARLOTTE ALBAEK
APPLICANT: WESTERGAARD, MAJKEN
APPLICANT: WESTERGAARD, MAJKEN
APPLICANT: WISSENBACH, MARGIT
TITLE OF INVENTION: OLIGOMERIC COMPOUNDS FOR THE MODULATION OF
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: 58614 (71.432)
CURRENT APPLICATION NUMBER: US/10/776,933
CURRENT APPLICATION NUMBER: US/10/776,933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/628,391
CURRENT FILING DATE: 2003-07-29
PRIOR APPLICATION NUMBER: PCT/IE02/00011
PRIOR FILING DATE: 2002-01-30
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: O'TOOLB, Dermot
APPLICANT: KELLEHOR, Dermot
APPLICANT: ABDEL-LATIF, Mohamed M.
TITLE OF INVENTION: INHIBITION OF NF-kappaB ACTIVATION
FILE REFERENCE: P69048US0
                                 PRIOR APPLICATION NUMBER: 60/446,374
PRIOR FILING DATE: 2003-02-10
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                   NUMBER OF SEQ ID NOS: 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE: OTHER INFORMATION: the redox active peptide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
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ORGANISM: Artificial sequence
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PatentIn Ver. 3.2
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Pred. No. 1
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APPLICANT: Waisman, David M.
APPLICANT: Kwon, Mijung
TITLE OF INVENTION: Anti-angiogenesis methods, con
FILE REFERENCE: ME02-001
CURRENT APPLICATION NUMBER: US/10/304,287
CURRENT FILING DATE: 2002-11-26
PRIOR APPLICATION NUMBER: US 60/333,866
PRIOR APPLICATION NUMBER: US 60/333,866
PRIOR FILING DATE: 2001-11-28
NUMBER OF SEQ ID NOS: 8
SOPTWARE: Microsoft Word
SEQ ID NO 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-290-072-233
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Matches
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APPLICANT: Heifetz, Peter
APPLICANT: Luginbuhl, Peter
APPLICANT: Luginbuhl, Peter
APPLICANT: Muchhal, Umesh
TITLE OF INVENTION: Nucleic Acids and Proteins with Thioredoxin Reductase Activity
FILE REFERENCE: A-71457-3
CURRENT FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: US/10/290,072
CURRENT FILING DATE: 2002-04-05
PRIOR FILING DATE: 2002-04-05
PRIOR APPLICATION NUMBER: US 60/376,682
PRIOR APPLICATION NUMBER: US 60/376,682
PRIOR APPLICATION NUMBER: US 10/141,531
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Pred. No. 1.7e+06;
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Publication No. US20040019104A1
GENERAL INFORMATION:
APPLICANT: Ebright, Richard H.
APPLICANT: Ebright, Yon W.
TITLE OF INVENTION: Reagents and Procedures
FILE REFERENCE: 744-34
          Sequence 2, Application US/10660118A
Publication No. US20040131606A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus
FILE REFERENCE: 2879-98
CURRENT APPLICATION NUMBER: US/10/660,118A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
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SEQ ID NO 5
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Best Local Similarity
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CURRENT FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: US 60/388,699
PRIOR FILING DATE: 2002-06-14
NUMBER OF SEQ ID NOS: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn version 3.2 
SEQ ID NO 233
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NAME/KEY: MISC FEATURE
(1)..(1)
PRIOR APPLICATION NUMBER: 60/462,082
                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
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OTHER INFORMATION: "Xaa" at position
OTHER INFORMATION: hydrophobic amino
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OTHER INFORMATION: "Xaa"
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OTHER INFORMATION: active site consensus sequence
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100.0%; Pred. No. 1.7e+06;
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Pred. No.
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PRIOR FILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.1
SEQ ID NO 2
LENGTH: 6
TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
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US-10-735-577-4
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US-10-660-118A-3
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                                                                                                                                            Sequence 4, Application US/10735577

Publication No. US20040142897A1

GENERAL INFORMATION:

APPLICANT: Waisman, David M.

TITLE OF INVENTION: Compositions and Methods

FILE REFERENCE: ME03-009
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SOFTWARE: Microsoft Word
SEQ ID NO 4
LENGTH: 6
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10 PRIOR APPLICATION NUMBER: 60/462,082 PRIOR FILING DATE: 2003-04-11 NUMBER OF SEQ ID NOS: 15 SOFTWARE: PatentIn version 3.1
                                                                     CURRENT APPLICATION NUMBER: US/10/735,577
CURRENT FILING DATE: 2003-12-12
PRIOR APPLICATION NUMBER: US 60/433,140
PRIOR FILING DATE: 2002-12-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Product and Process for Liquefaction FILE REFERENCE: 2879-98 CURRENT APPLICATION NUMBER: US/10/660,118A CURRENT FILING DATE: 2003-09-10
                                                 PRIOR FILING DATE: 2002-12-13
NUMBER OF SEQ ID NOS: 160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: White,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 6
TYPE: PAT
ORGANISM: Artificial sequence
FEATURE:
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NAME/KNY: misc feature
LOCATION: (1)...(6)
OTHER INFORMATION: Xaa = any amino acid
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Pred. No. 1.7
0; Mismatches
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                                                                 ; FEATURE:
; OTHER INFORMATION: C
US-10-954-951-11
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US-10-461-224-10
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Matches
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CURRENT APPLICATION NUMBER: US/10/461,224
CURRENT FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: US 60/388,699
PRIOR FILING DATE: 2002-06-14
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PATENTIN VERSION 3.2
SEQ ID NO 10
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Publication No. US20040019104A1
GENERAL INFORMATION:
APPLICANT: Ebright, Richard H.
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Best Local Similarity
   Query Match
Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                             SOFTWARE: PatentIn version 3.3 SEQ ID NO 11
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                                                                                                                                                                                                                                                                     APPLICANT: Hanson, George
APPLICANT: Kudlicki, Wieslaw Antoni
APPLICANT: Keppetipola, Shiranthi
TITLE OF INVENTION: Compositions and Methods For Synthesizing,
TITLE OF INVENTION: Detecting Biomolecules
FILE REFERENCE: 0942.6650001
                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/508,142
PRIOR FILING DATE: 2003-10-01
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/954,951
CURRENT FILING DATE: 2004-10-01
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TITLE OF INVENTION: Reagents and Procedures for High Specificity Labeling
                                                                                                                                  LENGTH: 8
TYPE: PRT
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Pred. No. 1.7e+06;
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Pred. No.
                  Score 31; I
Pred. No. 1
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nes 0;
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RESULT 17
US-10-615-343-2
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US-09-572-404B-970
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                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Complementary peptide ligands from the human genome FILE REFERENCE: Human patent CURRENT APPLICATION NUMBER: US/09/572,404B CURRENT FILING DATE: 2000-05-17 NUMBER OF SEQ ID NOS: 4203 SOFTWARE: ProtPatent version 1.0 SEQ ID NO 970 LENGTH: 10 TYPE: PRT
                 Sequence 2, Application US/10615343
Publication No. US20050009110A1
GENERAL INFORMATION:
APPLICANT: CHANG, XIAO-JIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 970, Applic Publication No. US20 GENERAL INFORMATION:
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APPLICANT: Proteom Ltd
APPLICANT: Proteom Ltd
TITLE OF INVENTION: Complementary peptide ligands from the human genome
FILE REFERENCE: Human patent
CURRENT APPLICATION NUMBER: US/09/572,404B
CURRENT FILING DATE: 2000-05-17
NUMBER OF SEQ ID NOS: 4203
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Best Local Similarity
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TITLE OF INVENTION: METHODS OF PRODUCING ANTIBODIES FOR DIAGNOSTICS AND THERAPEUTICS
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TYPE: PRT
                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: sequence located in LHB at 105-114 and may OTHER INFORMATION: this patent.
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4; Conservative 0; Mismatches 0
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nilarity 100.0%;
Conservative 0
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Pred. No. 2.1e+02;
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                                                                                                                                             US-10-044-844-217
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CURRENT APPLICATION NUMBER: US/10/615,343
CURRENT FILING DATE: 2003-07-08
NUMBER OF SEQ ID NOS: 22
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 217, Application US/10044844 Publication No. US20040009904A1 GENERAL INFORMATION:
                                                                       Matches
                                                                                      Query Match
Best Local Similarity
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Best Local
                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 40,130
REFERENCE/DOCKET NUMBER: 25231-20061.03
TELECOMPUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEFAX: (650) 494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO: 217:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH:
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Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/044,844
FILING DATE: 10-Jan-2002
CLASSIFICATION: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: CATHERIEE M. POLIZZI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 217:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER LLP
STREET: 755 PAGE MILL ROAD
CITY: PALO ALTO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: APL IMMUNOREACTIVE PEPTIDES, CONJUGATES THEREOF AND
METHODS OF TREATMENT FOR APL ANTIBODY-MEDIATED PATHOLOGIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Victoria, Edward J.
Marquis, David M.
Jones, David S.
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                                   2 CGPC 5
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                                                                     4;
 CGPC
                                                                                                                                                                                                                LENGTH: 16 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                       Conservative
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                                                                                     100.0%;
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                                                                     Score 31; DB 4; Pred. No. 2.9e+02; Mismatches 0;
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Pred. No. 2.6e+02;
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                                                                                                      Length 16;
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Sequence 10, Application US/09963339

Publication No. US20030049700A1

GENERAL INFORMATION:
APPLICANT: Bandaru, Rajasekhar
TITLE OF INVENTION: 22108 AND 47916, NOVEL HUMAN THIOREDOXIN
TITLE OF INVENTION: FAMILY MEMBERS AND USES THEREOF
FILE REFERENCE: 10448-090001
CURRENT APPLICATION NUMBER: US/09/963,339
CURRENT FILING DATE: 2001-09-25
PRIOR APPLICATION NUMBER: 60/235,049
PRIOR FILING DATE: 2000-09-25
INUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE: peptide
; SEQUENCE DESCRIPTION: SEQ ID NO: 217:
US-10-846-079-217
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GENERAL INFORMATION:
APPLICANT: Victo
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 217:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/846,079
FILING DATE: 13-May-2004
CLASSIFICATION: CUNknown>
ATTORNEY/ACENT INFORMATION:
NAME: CATHERINE M. POLIZZI
REGISTRATION NUMBER: 40,130
REFERENCE/DOCKET NUMBER: 25231-20061.03
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEPHONE: (650) 813-5600
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: APL IMMUNOREACTIVE PEPTIDES, CONJUGATES THEREOF AND METHODS OF TREATMENT FOR APL ANTIBODY-MEDIATED PATHOLOGIES
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TYPE: amino acid
STRANDEDNESS: single
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o. US20050208480A1
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Marquis, David M.
Jones, David S.
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100.0%; Pred. No. 2.9e+02;
htive 0; Mismatches 0;
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APPLICANT: Ruben et al.
APPLICANT: Ruben et al.
TITLE OF INVENTION: 27 Human secreted proteins
FILE REFERENCE: P2038P1
CURRENT APPLICATION NUMBER: U$/10/050,882
CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: 09/61,453
PRIOR FILING DATE: 2000-09-16
PRIOR APPLICATION NUMBER: PCT/U$00/06783
PRIOR APPLICATION NUMBER: 60/125,055
PRIOR FILING DATE: 1999-03-18
NUMBER OF SEQ ID NOS: 156
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 98
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; ORGANISM: Homo sapiens
US-10-145-586-59
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SEQ ID NO 59
LENGTH: 19
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Best Local Similarity
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Best Local Similarity
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TYPE: PRT
ORGANISM: Homo sapiens
-10-050-882-98
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CURRENT FILING DATE: 2002-05-14
Prior Application removed - See File Wrapp
NUMBER OF SEQ ID NOS: 95
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APPLICANT: Kapeller-Libermann, Rosana
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR FAMILY MEMBERS,
TITLE OF INVENTION: HUMAN THIOREDOXIN FAMILY MEMBERS, HUMAN LEUCINE-RICH
TITLE OF INVENTION: REPEAT FAMILY MEMBERS, AND HUMAN RINGFINGER FAMILY MEMBER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Alexandra Glucksmann, Maria
APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: M. Galvin, Katherine
APPLICANT: Weich, Nadine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 10448-188001
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TYPE: PRT
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No. US20030138890A1
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100.0%; Pred. No. 3.3e+02;
vative 0; Mismatches 0;
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Sequence 31, Application US/10179382

Publication No. US20030166235A1

GENERAL INFORMATION:
APPLICANT: THIREOS, GEORGE
APPLICANT: KAFETZOPOULOS, DIMITRIS
TITLE OF INVENTION: DNA ENCODING AN ARTHROPOD CHITIN SYNTHASE
FILE REFERENCE: DNA ENCODING AN ARTHROPOD CHITIN SYNTHASE
CURRENT APPLICATION NUMBER: US/10/179,382

CURRENT FILING DATE: 2002-06-26
PRIOR APPLICATION NUMBER: US/09/230,041

PRIOR APPLICATION SYNTH

PRIOR APPLICATION NUMBER: US/09/230,041

PRIOR EILING DATE: 1999-01-19

NUMBER OF SEQ ID NOS: 42
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FILE REFERENCE: PZ038P1C2
CURRENT APPLICATION NUMBER: US/10/963,903
CURRENT FILING DATE: 2004-10-14
PRIOR FILING DATE: 2004-10-14
PRIOR APPLICATION NUMBER: US 10/050,882
PRIOR FILING DATE: 2002-01-18
PRIOR FILING DATE: 2000-09-13
PRIOR FILING DATE: 2000-09-13
PRIOR FILING DATE: 2000-09-13
PRIOR FILING DATE: 2000-03-16
PRIOR FILING DATE: 108-03-18
PRIOR FILING DATE: 108-03-18
PRIOR FILING DATE: 1099-03-18
PRIOR FILING DATE: 1099-03-18
PRIOR FILING DATE: 1099-03-18
NUMBER OF SEQ ID NOS: 163
SOFTWARE: Patentin Ver: 2.0
                                                                                                                                 ; TYPE: PRT
; ORGANISM: Rhizobium leguminosarum
US-10-179-382-31
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US-10-179-382-31
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US-10-963-903-98
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                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 31
LENGTH: 28
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LENGTH: 25
TYPE: PRT
ORGANISM: Homo sapiens
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Publication No. US20050239099A1
GENERAL INFORMATION:
                                              Query Match
Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 4; Conserv
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      2 CGPC 5
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                                                   Conservative
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                                              100.0%; Score 31; DB 4;
100.0%; Pred. No. 4.2e+02;
tive 0; Mismatches 0;
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                                                                                    Length 28;
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21 CGPC 24

FOR

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APPLICANT: Bupp, Keith;
APPLICANT: University of Medicine and Dentistry of New Jersey;
APPLICANT: University of Medicine and Dentistry of New Jersey;
TITLS OF INVENTION: Targeting Viral Vectors to Specific Cells;
FILE REFERENCE: 601-1-095
CURRENT APPLICATION NUMBER: US/09/881,572A
CURRENT FILING DATE: 2002-03-05
PRIOR APPLICATION NUMBER: 60/212,239
PRIOR APPLICATION NUMBER: 60/212,239
PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.0
SEQ ID NO 5
LENGTH: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Amphotropic Murine Leukemia Virus US-09-881-572A-25
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US-09-864-761-40069
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                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: PRT; ORGANISM: amphotropic US-09-881-572A-5
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US-09-881-572A-5
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SOFTWARE: Patentin version 3.0
SEQ ID NO 25
LENGTH: 29
Sequence 40069, Application US/09864761
Patent No. US20020048763A1
GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
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APPLICANT: Roth, Monica;
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Patent No. US20020164583A1
                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 4; Conserv
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Best Local Similarity 100.0%;
Matches 4; Conservative 0
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APPLICANT: University of Medicine and Dentistry of New Jersey
TITLE OF INVENTION: Targeting Viral Vectors to Specific Cells
FILE REFERENCE: 601-1-095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/881,572A
CURRENT FILING DATE: 2002-03-05
PRIOR APPLICATION NUMBER: 60/212,239
PRIOR FILING DATE: 2000-06-16
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Pred. No. 4.4e+02;
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Pred. No.
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PRIOR FILING DATE: 2000-10-04
PRIOR PELICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-08-03
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SEQ ID NO 40069
                                                                                                                                  Query Match
Best Local &
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NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers.
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2000-06-30 PRIOR APPLICATION NUMBER: US 09/774,203
                                                                                                                                                                                                                                     OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 5.4
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 4.6
OTHER INFORMATION: EXPRESSED IN HETAL LIVER, SIGNAL = 4.4
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 4.9
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 4.3
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 4.3
OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 4.3
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 5
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 4
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 4
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 4
OTHER INFORMATION: SWISSPROT HIT: BF446535.1, EVALUE 6.00e-12
OTHER INFORMATION: SWISSPROT HIT: Q02597, EVALUE 1.70e+00
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TYPE: PRT
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OTHER INFORMATION: EXPRESSED IN LUNG
OTHER INFORMATION: EXPRESSED IN HELA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 2000-09-21
APPLICATION NUMBER: US 09/608,408
FILING DATE: 2000-06-30
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APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00663
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                                                     CGPC 5
CGPC 11
                                                                                                           Conservative
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                                                                                                      100.0%; Sc
100.0%; P:
ative 0;
                                                                                                        Score 31; DB
Pred. No. 4.6
0; Mismatches
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                                                                                                                                                            Length 32
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RESULT 28 US-10-424-599-160512

Sequence 160512, Application US/10424599

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RESULT 30
US-09-801-260-5
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FILE REFERENCE: BB1367 US NA
CURRENT APPLICATION NUMBER: US/10/044,359
CURRENT FILING DATE: 2002-01-11
PRIOR APPLICATION NUMBER: 09/599,416
PRIOR APPLICATION NUMBER: 69/140,227
PRIOR APPLICATION NUMBER: 60/140,227
PRIOR FILING DATE: 1999-06-22
PRIOR FILING DATE: 1999-06-22
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US-10-424-599-160512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Androctonus mauretanicus US-10-044-359-26
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Best Local S
Matches 4
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APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Cao Yinua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 160512
Sequence 5, Application US/09801260
Patent No. US20020034801A1
GENERAL INFORMATION:
APPLICANT: Curtis, Rory A.J.
TITLE OF INVENTION: 22105, A NOVEL HUMAN THIOREDOXIN FAMILY
TITLE OF INVENTION: MEMBER AND USES THEREOF
FILE REFERENCE: 10448-022001
CURRENT APPLICATION NUMBER: US/09/801,260
CURRENT FILING DATE: 2001-03-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Herrman, Rafael
APPLICANT: Wong, James F.
APPLICANT: Lee, Jian-Ming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Microsoft Office 97
SEQ ID NO 26
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Glycine max
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PRIOR APPLICATION NUMBER: US 60/187,447

PRIOR FILING DATE: 2000-03-07

NUMBER-OF SEQ ID NOS: 5

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 5

LENGTH: 36

TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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US-10-145-586-44
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                                                                                                                                                                                                                                                       US-10-450-763-51435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-145-586-44
                                                                                                                                      Sequence 51435, Application US/10450763
Publication No. US20050196754A1
GENERAL INFORMATION:
APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 44, Application US/10145586 Publication No. US20030138890A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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SEQ ID NO 44
LENGTH: 36
TYPE: PRT
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FILE REFERENCE: 790CIP3/US
CURRENT APPLICATION NUMBER: US/10/450,763
CURRENT FILING DATE: 2003-06-11
PRIOR APPLICATION NUMBER: PCT/US01/08631
PRIOR FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: 09/540,217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/145,586
CURRENT FILING DATE: 2002-05-14
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 95
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APPLICANT: Bandaru, Rajasekhar
APPLICANT: Bandaru, Rajasekhar
APPLICANT: Kappeller-Libermann, Rosana
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR FAMILY MEMBERS,
TITLE OF INVENTION: HUMAN THIOREDONIN FAMILY MEMBERS, HUMAN LEUCINE-RICH
TITLE OF INVENTION: REPEAT FAMILY MEMBERS, AND HUMAN RINGFINGER FAMILY MEMBER
TITLE OF INVENTION: REPEAT FAMILY MEMBERS, AND HUMAN RINGFINGER FAMILY MEMBER
FILE REFERENCE: 10448-18801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: M. Galvin, Ka
APPLICANT: Weich, Nadine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Alexandra Glucksmann, Maria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
OTHER INFORMATION: consensus sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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100.0%; Pred. No. 5e-
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 31; DB 4; 100.0%; Pred. No. 5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Katherine
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PRIOR FILING DATE: 2000-03-31

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US-09-864-761-35367; Sequence 35367, Application US/09864761; Patent No. US20020048763A1
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                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 96/22,366
PRIOR APPLICATION NUMBER: US 96/32,366
PRIOR APPLICATION NUMBER: US 96/32,366
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR PRILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR PILING DATE: 2001-01-30
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PRIOR FILING DATE: 2000-08-23
NUMBER OF SEQ ID NOS: 60736
SOFTWARE: CUSECOM
SEQ ID NO 51435
                                                                               PRIOR FILING DATE:
PRIOR APPLICATION NI
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PRIOR FILING DATE:
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Best Local Similarity
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence
SEQ ID NO 35367
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: Aeomica-X-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
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                                                                                                                    APPLICATION NUMBER: US 09/774,203
                                                                                                                                                                                                                   FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
                                                                                                                                             FILING DATE: 2000-09-21
APPLICATION NUMBER: US 09/608,408
FILING DATE: 2000-06-30
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l, David K.
                                                                                     2001-01-29
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Pred. No.
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US-10-424-599-255327; Reguence 255327, Application US/10424599; Publication No. US20040031072A1
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US-10-293-157-20
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Matches
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Publication No. US20030144200A1
GENERAL INFORMATION:
APPLICANT: BAIRD, ANDREW
APPLICANT: ANDREASON, GRAI
                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local 9
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: NOVEL FORMS OF THE ANGIOGENIC FACTOR TITLE OF INVENTION: VASCULAR ENDOTHELIAL CELL GROWTH FACTOR FILE REFERENCE: 240/086
              APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zbou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US/09/244,583
PRIOR FILING DATE: 1999-02-04
PRIOR APPLICATION NUMBER: 60/073,979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/293,157
CURRENT FILING DATE: 2002-11-12
CURRENT APPLICATION NUMBER: US/10/424,599
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ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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R INFORMATION:
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NN: EXPRESSED IN LUNG, SIGNAL = 1.5

NN: EXPRESSED IN BRAIN, SIGNAL = 2.7

NN: EXPRESSED IN BRAIN, SIGNAL = 2.7

NN: EXPRESSED IN HOME MARROW, SIGNAL = 3.4

NN: EXPRESSED IN ADULT LIVER, SIGNAL = 3.4

NN: EXPRESSED IN HOLLT LIVER, SIGNAL = 2.4

NN: EXPRESSED IN HELLA, SIGNAL = 2.2

NN: EXPRESSED IN HELLA, SIGNAL = 1.8

NN: EXPRESSED IN HELLAOO, SIGNAL = 1.8

NN: EXPRESSED IN HELART, SIGNAL = 3.2

NN: EXPRESSED IN HELART, SIGNAL = 3.0

NN: EXPRESSED IN HELART, SIGNAL = 3.0

NN: EXPRESSED IN HELART, SIGNAL = 3.00e-03

NN: EXPRESSED IN HIT: AW665671.1, EVALUE 3.00e-04
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                                                                                                                                                                                                                                                                                                                                           Score 31; DE
Pred. No. 5.7
0; Mismatches
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Pred. No.
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NUMBER OF SEQ II SEQ ID NO 255327

ID NOS:

2003-04-28 285684

LENGTH:

CURRENT FILING DATE:

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; TYPE: PRT; ORCANISM: Glycine max; PEATURE: FEATURE: PAT_MRT3847_72582C.1.pep US-10-424-599-255327
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Homo sapiens US-09-832-355A-3
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                                               CURRENT APPLICATION NUMBER: US/10/104,440
CURRENT FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: 09/580,803
PRIOR FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: 60/069,155
PRIOR FILING DATE: 1997-12-09
PRIOR APPLICATION NUMBER: 60/069,687
PRIOR APPLICATION NUMBER: 60/069,687
PRIOR APPLICATION NUMBER: 60/069,587
PRIOR APPLICATION NUMBER: 60/078,541
PRIOR FILING DATE: 1998-03-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.
"Arches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.
Warches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 3, Application US/09832355A Publication No. US20030027751A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.0 SEQ ID NO 3
SEQ ID NO 11
                                                                                                                                                                                                                                                                                                                                     Sequence 11, Applica Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Kovesdi, Imre
APPLICANT: Kessler, Paul
TITLE OF INVENTION: VEGF FUSION PROTEINS
FILE REFERENCE: 205654
CURRENT APPLICATION NUMBER: US/09/832,355A
CURRENT FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 126
                                                                                                                                                                                                                                         TITLE OF INVENTION: ANTAGONISTS OF NEUROPILIN RECEPTOR FUNCTION AND TITLE OF INVENTION: THEREOF
                NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                          TITLE OF INVENTION: THEREOF
FILE REFERENCE: 48802 C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 44
TYPE: PRT
                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                  APPLICANT: KLAGSBRUN, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18 CGPC 21
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                                                                                                                                                                                                                                                                                 SOKER, She
                                                                                                                                                                                                                                                                                                                                                   Application US/10104440 No. US20020132774A1
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100.0%; Pred. No. 5.
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Pred. No. 5.7
0; Mismatches
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APPLICANT: KLAGSBRUN, Michael
APPLICANT: SOKER, Shay
APPLICANT: SOKER, Shay
APPLICANT: GAGNON, Michael L.
TITLE OF INVENTION: SOLUBLE INHIBITORS OF VASCULAR ENDOTHELIAL GROWTH
TITLE OF INVENTION: ADD USE THEREOF
FILE REFERENCE: 48801 C
CURRENT PILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: US/10/104,610
CURRENT FILING DATE: 2000-05-30
PRIOR FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: 60/69,155
PRIOR APPLICATION NUMBER: 60/69,155
PRIOR APPLICATION NUMBER: 60/69,155
PRIOR APPLICATION NUMBER: 60/69,155
PRIOR APPLICATION NUMBER: 60/69,157
PRIOR APPLICATION NUMBER: 60/69,687
PRIOR APPLICATION NUMBER: 60/069,687
PRIOR APPLICATION NUMBER: 60/069,687
PRIOR FILING DATE: 1997-12-12
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US-10-104-440-11
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US-10-424-599-252212
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US-10-104-610-15
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Best Local Similarity 100.
Best Local Similarity 200.
                                                                                                                                                    APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285884
SEQ ID NO 252212
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 252212, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT:
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APPLICANT: Kovalic David K
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TYPE: PRT
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                                                        NAME/KEY: unsure
                                                                                                                             LENGTH: 45
TYPE: PRT
               LOCATION: (1)..(45)
OTHER INFORMATION: unsure at
                                                                                                  ORGANISM: Glycine max
EATURE:
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                  all Xaa locations
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                                                 ; OTHER INFORMATION: Clone ID: MRT4577_56384C.1.pep US-10-425-115-321622
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APPLICANT: Kovalic, David K.
APPLICANT: Chou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21 (53222)B
CURRENT EAPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 210616
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Best Local Similarity 100
Matches 4; Conservative
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
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Best Local Similarity
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SEQ ID NO 321622
LENGTH: 47
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                       APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21 (53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Zea mays
                                                                               FEATURE: Unsure
LOCATION: (1)..(47)
OTHER INFORMATION: unsure at
FEATURE:
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100.0%;
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                                                                                                      all Xaa locations
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Pred. No. 5.
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Pred. No. 6e+02;
 Score 31;
Pred. No.
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 DB 4;
6e+02;
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                 Length 47;
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CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: 60/157,832
PRIOR FILING DATE: 1999-10-05
PRIOR PPLICATION NUMBER: 60/160,191
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR FILING DATE: 1999-10-28
PRIOR PILING DATE: 1999-10-28
PRIOR PILING DATE: 1999-10-18
PRIOR APPLICATION NUMBER: 60/164,769
PRIOR APPLICATION NUMBER: 60/164,769
PRIOR FILING DATE: 1999-11-12
PRIOR FILING DATE: 1999-11-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/173,383
PRIOR FILING DATE: 1999-12-28
PRIOR APPLICATION NUMBER: 60/175,693
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/184,831
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: 60/191,637
PRIOR FILING DATE: 2000-03-23
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                                                                                                                                                                                                                                                                                                                                  Sequence 334172, Application US/10425115
Publication No. US20040214272A1
GENERAL INFORMATION:
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                                                                                 CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 334172
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Best Local Similarity
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                                                                                                                                                                      APPLICANT: La ROSA, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
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APPLICANT: et al.
TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: DROSOPHILA GENES.
TITLE OF INVENTION: DROSOPHILA GENES.
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                    LENGTH: 49
TYPE: PRT
ORGANISM: Zea mays
FEATURE:
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Pred. No. 6e+02;
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 52
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APPLICANT: CUITIS, ROLY A.J.
TITLE OF INVENTION: 22105, A NOVEL HUMAN THIOREDOXIN FAMILY
TITLE OF INVENTION: MEMBER AND USES THEREOF
FILE REFERENCE: 10448-022001
                                                            SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 43
                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Alexandra Glucksmann, Maria
APPLICANT: Silos-Santiago, Inmaculada
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 43, Application US/10145586 Publication No. US20030138890A1
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Best Local Similarity
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                                                                                                                                                                                                      APPLICANT: Bandaru, Rajasekhar
APPLICANT: Kapeller-Libermani, Rosana
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR FAMILY MEMBERS,
TITLE OF INVENTION: HUMAN THIOREDOXIN FAMILY MEMBERS, HUMAN LEUCINE-RICH
TITLE OF INVENTION: REPEAT FAMILY MEMBERS, AND HUMAN RINGFINGER FAMILY MEMBER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/801,260
CURRENT FILING DATE: 2001-03-06
PRIOR APPLICATION NUMBER: US 60/187,447
PRIOR FILING DATE: 2000-03-07
                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/145,586
CURRENT FILING DATE: 2002-05-14
                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                     Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 95
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OTHER INFORMATION: unsure at all Xaa locations
FEATURE:
                       TYPE: PRT
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                       REFERENCE: 10448-188001
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                                                                                                                                                                                                                                                                                                                                                     M. Galvin, Katherine
                                                                                                                                                                                                                                                                                                                                    Weich, Nadine
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Pred. No. 6.2e+02;
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US-10-424-599-166951
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US-10-437-963-164055
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                                 US-10-424-599-166951
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                                                                                                                             APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5223)B
CURRENT PELICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 16951
                                                                                                                                                                                                                                                                                     Sequence 166951, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
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SEQ ID NO 164055
LENGTH: 52
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APPLICANT: La Rosa, Thomas
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Best Local (
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Best Local Similarity
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APPLICANT:
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CURRENT FILING DATE: 2003-05-14
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                                          FEATURE: OTHER INFORMATION: Clone ID: PAT_MRT3847_121771C.1.pep
                                                                             TYPE: PRT ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Oryza sativa FEATURE:
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                                                                                                                    LENGTH: 53
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Wu, Wei
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Kovalic, David K.
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CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
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Matches 4; Conserv
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                                            OTHER INFORMATION: NOTHER INFORMATION: N
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APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
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                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
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R APPLICATION N
R FILING DATE:
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APPLICATION NUMBER: PCT/US01/00668
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Hanzel, David K.
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                                        MAP TO ACO11510.3
EXPRESSED IN BRAIN
EXPRESSED IN ADULT
EXPRESSED IN HEART
EXPRESSED IN LUNG,
EXPRESSED IN HELACE
EXPRESSED IN HELACE
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DD IN BRAIN, SIGNAL = 2.00 IN ADULT LIVER, SIGNAL = 1.00 IN HEART, SIGNAL = 1.00 IN LUNG, SIGNAL = 2.4 IN LACENTA, SIGNAL = 1.7 IN HELA, SIGNAL = 2.4 IN HELA, SIGNAL = 2.4 IN HELA, SIGNAL = 2.4 IN FETAL LIVER, SIGNAL E 
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; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.7; OTHER INFORMATION: EST HUMAN HIT: BE148597.1, EVALUE 2.00e-24; OTHER INFORMATION: SWISSPROT HIT: P03204, EVALUE 3.50e+00 US-09-864-761-42458
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                                               US-10-425-115-307743
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APPLICANT: La Rosa, Thomas
APPLICANT: Kovalic, David
APPLICANT: Xovalic, Yihua
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 10011-00001
CURRENT APPLICATION NUMBER: US/10/318,302
CURRENT FILING DATE: 2002-12-12
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin version 3.1
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: POSTES
APPLICANT: Chae,
APPLICANT: Chae,
APPLICANT: Gho,
APPLICANT: Yang,
APPLICANT: Kwon,
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Publication No. US20030171556A1
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Best Local (
                                                                                                                                       NUMBER OF SEQ ID NOS:
SEQ ID NO 307743
                                                                                                                                                                                                                                                                                                                                Sequence 307743, Application US/10425115 Publication No. US20040214272A1
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              Query Match
Best Local Similarity
                                                                                                                                                                   APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules .
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Bae, Dong-Goo
APPLICANT: Hwang, Sewook
TITLE OF INVENTION: BETA-AMYLOID BINDING FACTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 55
TYPE: PRT
ORGANISM: Homo sapiens
                                                                           TYPE: PRT
ORGANISM: Zea mays
FEATURE:
                                                          OTHER INFORMATION: Clone ID: MRT4577_43730C.1.pep
                                                                                                                          ENGTH: 55
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4; Conservative
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Kwon, Byung Oh
Bae, Dong-Goo
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31;
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. 6.7e+02;
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              DB 4;
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-868-577A-28
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 28
LENGTH: 55
                                                               NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 26469
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: DETECTION KIT, SUCH AS TITLE OF INVENTION: ARRAYS, FOR DETECTING TITLE OF INVENTION: DROSOPHILA GENES.
                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/173,383
PRIOR FILING DATE: 1999-12-28
PRIOR APPLICATION NUMBER: 60/175,693
PRIOR FILING DATE: 2000-01-12
                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/161,932 PRIOR FILING DATE: 1999-10-28
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CURRENT FILING DATE: 2005-04-04
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APPLICANT: et al.
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PRIOR FILING DATE: 2003-09-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/157,832
ORGANISM: DROSOPHILA
                                                                                                                                                         APPLICATION NUMBER: 60/191,637
                                                                                                                                                                                                       APPLICATION NUMBER: 60/184,831
                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 60/164,769
FILING DATE: 1999-11-12
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Pred. No. 6.7e+02;
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EXPRESSION OF 10,000 OR
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APPLICANT: et al.

APPLICANT: et al.

TITLE OF INVENTION: DETECTION KIT, SUCH AS NU
TITLE OF INVENTION: ARRAYS, FOR DETECTING EX:
TITLE OF INVENTION: DROSOPHILA GENES.

FILE REFERENCE: CL000728

CURRENT APPLICATION NUMBER: US/11/097,143

CURRENT APPLICATION NUMBER: 60/157,832

PRIOR APPLICATION NUMBER: 60/160,191

PRIOR APPLICATION NUMBER: 60/160,191

PRIOR FILING DATE: 1999-10-19

PRIOR APPLICATION NUMBER: 60/161,932

PRIOR FILING DATE: 1999-10-28

PRIOR APPLICATION NUMBER: 60/161,932

PRIOR APPLICATION NUMBER: 60/164,769

PRIOR APPLICATION NUMBER: 60/164,769

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PRIOR APPLICATION NUMBER: 60/164,769
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US-11-097-143-26490
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US-10-425-115-219153
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APPLICANT: LA ROSA, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yongwei
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Best Local S
Matches 4
                                                               PRIOR FILING DATE: 1999-11-12
PRIOR APPLICATION NUMBER: 60/173,383
PRIOR FILING DATE: 1999-12-28
PRIOR APPLICATION NUMBER: 60/175,693
PRIOR PILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/184,831
PRIOR PILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 26490, Application US/11097143 Publication No. US20050208558A1 GENERAL INFORMATION:
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LENGTH: 56
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TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT APPLICATION OND NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
                     PRIOR APPLICATION NUMBER: 60/191,637
PRIOR FILING DATE: 2000-03-23
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ORGANISM: Zea mays
FEATURE:
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les 4; Conserv
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100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0;
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EXPRESSION OF 10,000 OR MORE
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                                                                                                                                                                                                                                                                                                    RESULT 56
US-10-425-115-353222
US-10-425-115-353222, Application US/10425115
; Publication No. US/0040214272A1
; GENERAL INFORMATION:
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APPLICANT: Rank, David K.
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
FILE REFERENCE: AEOMICA-X-2
CURRENT APPLICATION NUMBER: US/10/029,386
CURRENT FILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 28568
                                   APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yongwei
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REPERENCE: 38-21(5)222)B
FILE REPERENCE: 38-21(5)222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 353222
LENGTH: 57
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
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Publication No. US20030194704A1
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Best Local (
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OTHER INFORMATION: MAP TO CHR2.1

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.5

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.71

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 2.71

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.7

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.3

OTHER INFORMATION: SWISSPROT HIT: P00203, EVALUE 1.20e+00
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ORGANISM: Zea mays
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Similarity 100.0%;
4; Conservative 0.
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Pred. No. 6.8e+02;
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Pred. No. 6.9e+02;
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US-10-424-599-185374
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                                     ; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_97667C.1.pep
US-10-437-963-202381
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                                                                                                                                                   TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILLING DATE: 2003-65-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 202381
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SEQ ID NO 185374
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
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Publication No. US20040031072A1
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Best Local Similarity
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APPLICANT:
APPLICANT:
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)8
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
Query Match
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ORGANISM: Glycine max
                                                                                             TYPE: PRT
ORGANISM: Oryza sativa
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Wu, Wei
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Barbazuk, Brad
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100.0%; Pred. No. 7e-
tive 0; Mismatches
   Score 31;
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   DB 4;
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   Length 58;
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RESULT 61
US-11-097-143-10200
; Sequence 10200, Application US/11097143
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US-09-864-408A-7494
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Best Local Similarity
Matches 4; Conserva
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; OTHER INFORMATION: Clone ID: PAT_MRT3847_130272C.1.pep
US-10-424-599-176362
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US-10-424-599-176362
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7494
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Publication No. US20040031072A1
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TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/864,408A CURRENT FILING DATE: 2001-05-24 PRIOR APPLICATION NUMBER: 60/206,690
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: No. US20040009474A1el Human Polynucleotides and Polypeptides Enc.
FILE REFERENCE: 21402-012
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
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nes 4; Conserv
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Pred. No.
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; OTHER INFORMATION: Clone ID: PAT_MRT3847_93452C.1.pep US-10-424-599-278438
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US-11-097-143-10200
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                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 278438
                                                                                         Best Loc
Matches
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APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 43008
SOPTWARE: FASTSEQ for Windows Version
SEQ ID NO 10200
LENGTH: 60
TYPE: PRT
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APPLICANT: Venter, J. Craig
APPLICANT: et al.
TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
TITLE OF INVENTION: DROSOPHILA GENES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 278438, Application US/10424599 Publication No. US20040031072A1
                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: 60/191,637
PRIOR FILING DATE: 2000-03-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1999-11-12
PRIOR APPLICATION NUMBER: 60/173,383
PRIOR FILING DATE: 1999-12-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 1999-10-19
PRIOR APPLICATION NUMBER: 60/161,932
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/11/097,143
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: 60/157,832
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: 60/160,191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/184,831
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/164,769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 1999-10-28
                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                  ORGANISM: Glycine max
                                                                                                                                                                                                                                                                             TYPE: PRT
                                                                                                                                                                                                                                                                                                LENGTH: 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       / Match 100.0%; Score 31; DB 6; 1
Local Similarity 100.0%; Pred. No. 7.1e+02;
1es 4; Conservative 0; Mismatches 0;
                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41 CGPC 44
28 CGPC 31
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                                              2 CGPC 5
                                                                                         4
                                                                                       Conservative
                                                                                                             100.0%; Score 31; DB 4; 100.0%; Pred. No. 7.3e+02;
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                                                                                            Mismatches
                                                                                                                                     Length 62;
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                                                                                            Indels
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US-10-425-115-233772

Sequence 233772, Application US/10425115

Publication No. US20040214272A1

GENERAL INFORMATION:

APPLICANT: La Rosa, Thomas J.

APPLICANT: Kovalic, David K.

APPLICANT: Cao, Yihua

APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and TITLE OF INVENTION: Plants

TITLE OF INVENTION: Plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 26472, Application US/11097143

Publication No. US20050208558A1

GENERAL INFORMATION:

APPLICANT: Venter, J. Craig

APPLICANT: venter, J. Craig

APPLICANT: et al.

TITLE OF INVENTION: DETECTION KIT, SUCH AS NU
TITLE OF INVENTION: DROSOPHILA GENES.

FILE REFERENCE: CL000728

FILE REFERENCE: CL000728
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Best Local Similarity
""" hes 4; Conservi
                                                                    LENGTH: 63
TYPE: PRT
ORGANISM: DROSOPHILA
US-11-097-143-26472
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US-11-097-143-26472
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; OTHER INFORMATION: Clone
US-10-425-115-233772
                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/11/097,143
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: 60/157,832
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: 60/160,191
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR APPLICATION NUMBER: 60/164,769
PRIOR APPLICATION NUMBER: 60/164,769
PRIOR APPLICATION NUMBER: 60/164,769
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SOFTWARE: FastSEQ for
SEQ ID NO 26472
LENGTH: 63
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LENGTH: 63
Query Match
Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/191,637 PRIOR FILING DATE: 2000-03-23
                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/184,831 PRIOR FILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/173,383
PRIOR PELLING DATE: 1999-12-8
PRIOR APPLICATION NUMBER: 6-28
PRIOR FILING DATE: 2000-01-12
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ORGANISM: Zea mays
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                                                                                                                                                                                      for Windows Version
100.0%;
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Score 31; DB 6;
Pred. No. 7.4e+02;
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EXPRESSION OF 10,000 OR MORE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and Other Molecules Associated With
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                                                                               US-10-424-599-180203
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CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 336734
                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 180203
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Best Local Similarity
Matches 4; Conserv
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Query Match
Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 64
TYPE: PRT
ORGANISM: Zea mays
FEATURE:
                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53223)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants FILE REFERENCE: 38-21(53222)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Kovalic, David
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: La Rosa, Thomas J. APPLICANT: Kovalic, David K.
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APPLICANT: Cao Yongwei
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LOCATION: (1)..(64)
OTHER INFORMATION: unsure at all Xaa locations
                                                                                                                           FEATURE:
NAME/KEY: unsure
LOCATION: (1).. (65)
OTHER INFORMATION: unsure at all Xaa locations
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                                                                                                                                                                                                                                         LENGTH: 6
TYPE: PRT
                                                                                               FEATURE:
OTHER INFORMATION: Clone ID:
                                                                                                                                                                                                                 ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   40 CGPC 43
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100.0%; Score 31; DB 4; ilarity 100.0%; Pred. No. 7.5e+02; Conservative 0; Mismatches 0;
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                                      Length 65;
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US-10-425-115-256158
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                                                                           Query Match
Best Local S
Matches 4
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SEQ ID NO 256158
LENGTH: 67
TYPE: PRT
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
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LENGTH: 67
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                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleac Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
FULL REFERENCE: 38-21(53221)B
FULL REFERENCE: US/10/437,963
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILLING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
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                                                                                                                                                                                                              ORGANISM: Zea mays
                                                                                                                                                                       OTHER INFORMATION: Clone ID: MRT4577_165207C.1.pep
                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     y Match 100.0%;
Local Similarity 100.0%;
hes 4; Conservative 0;
                                                                                                Local Similarity
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10 CGPC 13
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Yihua
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Wu, Wei
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Barbazuk, Brad
                                                                             Conservative
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                                                                                              100.0%; Score 31; DB 4; 100.0%; Pred. No. 7.7e+02;
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US-11-097-143-32565
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                                                               FILE REFERENCE: 807A
CURRENT APPLICATION NUMBER: US/10/243,552
CURRENT APPLICATION NUMBER: US 60/322,511
PRIOR APPLICATION NUMBER: US 60/322,511
PRIOR FILING DATE: 2001-09-13
PRIOR PPLICATION NUMBER: PCT/US00/35017
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: US 09/488,725
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PRIOR APPLICATION NUMBER: US 09/488,725
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                                                                                                                                                                                                                                                                                                                                         APPLICANT: Yang, Yonghong
APPLICANT: Wang, Zhiwei
APPLICANT: Weng, Gzhi
APPLICANT: Weng, Gzhi
APPLICANT: Meng, Gzhi
TITLE OF INVENTION: Novel Nucleic Acids and
TITLE OF INVENTION: Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: CL000728
CURRENT APPLICATION NUMBER: US/11/097,143
CURRENT FILING DATE: 2005-04-04
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TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
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PRIOR FILING DATE: 2000-03-23
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PRIOR FILING DATE: 1999-10-19
PRIOR APPLICATION NUMBER: 60/161,932
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 2000-02-24
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PRIOR FILING DATE: 2000-01-12
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PRIOR APPLICATION NUMBER: 60/164,769
PRIOR FILING DATE: 1999-11-12
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APPLICATION NUMBER: US 09/491,404
                  APPLICATION NUMBER: PCT/US01/02623 FILING DATE: 2001-01-25
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Pred. No. 7.7e+02;
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PRIOR APPLICATION NUMBER: US 09/496,914
PRIOR APPLICATION NUMBER: US 09/496,914
PRIOR PILING DATE: 2000-02-03
PRIOR APPLICATION NUMBER: US 09/560,875
PRIOR PILING DATE: 2000-04-27
PRIOR FILING DATE: 2000-04-27
PRIOR PILING DATE: 2000-04-27
PRIOR PILING DATE: 2001-02-26
Remaining Prior Application data removed - See File Wrapper or PALM.
VUMBER OF SEQ ID NOS: 998
SOFTWARE: Pt FL_genes Version 5.0
SEQ ID NO 978
SEQ ID NO 978
PTYPE: PRT
ORGANISM: Homo sapiens
US-10-243-552-978

Query Match
Best Local Similarity 100.0%; Score 31; DB 4; Length 68;
Best Local Similarity 100.0%; Pred. No. 7.8e+02;
Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 2 CGPC 5
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Search completed: February 23, 2006, 00:47:46
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GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use There
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/560,118
PRIOR PILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/550,865
PRIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin version 3.3
SEQ ID NO 1
LENGTH: 4
TYDE: PRT
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US-11-090-916-1
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Publication No. US20050260140A1
GENERAL INFORMATION:
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Best Local
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US-11-183-136-38
US-11-072-512-2161
US-11-098-686-11331
US-11-098-398-40
US-10-980-388-40
US-10-980-388-40
US-11-022-289-7
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US-11-0453-372-236
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Pred. No. 8.2e+04;
; Mismatches 0;
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11-077-386-25
11-183-136-38
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RESULT 4
US-11-23-547-1
; Sequence 1, Application US/11223547
; Publication No. US20060009527A1
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RESULT

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GENERAL INFORMATION:
APPLICANT: Richon, Victoria M.
APPLICANT: Richon, Victoria M.
APPLICANT: Rifkind, Richard A.
APPLICANT: Butler, Lisa M.
TITLE OF INVENTION: Method of Treating TRX Mediated Dis
FILE REFERENCE: 3254.1005-001
CURRENT APPLICATION NUMBER: US/11/223,405
CURRENT FILING DATE: 2005-09-09
PRIOR APPLICATION NUMBER: US/10/369,094
PRIOR FILING DATE: 2003-02-14
PRIOR FILING DATE: 2003-02-14
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CURRENT FILING DATE: 2005-06-03
PRIOR APPLICATION NUMBER: US 60/577,089
PRIOR FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 60/357,383
PRIOR FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin version 3.3
SEQ ID NO 8
LENGTH: 4
TYPE: PRT
CORONNESS: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Artificial
; FEATURE:
; OTHER INFORMATION: Synthetic peptide
US-11-144-301A-8
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Best Local Similarity luv.
"---hes 4; Conservative
                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: U.S. 60/357,383
PRIOR FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FASUSEQ for Windows Version 4.0
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 8, Application US/11144301A Publication No. US20050288227A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/11223405 Publication No. US20060009526A1
                                                                                 Query Match 100.0%; Score 31;
Best Local Similarity 100.0%; Pred. No.
Matches 4; Conservative 0; Mismatch
                                                                                   Matches
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APPLICANT: Johanna, Ungerstedt
TITLE OF INVENTION: USE OF THIOREDOXIN MEASUREMENTS
FILE REFERENCE: 28517-501 UTIL
                                                                                                                                                                                                                LENGTH: 4
TYPE: PRT
                                                                                                                                                                                          ORGANISM: homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 CGPC 5
                                         2 CGPC
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100.0%; Pred. No. 8.3
tive 0; Mismatches
                                                                                     Mismatches
                                                                                                         8.2e+04;
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                                                                                                                                 DB 7;
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GENERAL INFORMATION:

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                                                                       Query Match
Best Local S
Matches 4
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/11/223,547
CURRENT FILING DATE: 2005-09-09
PRIOR APPLICATION NUMBER: US/10/369,094
PRIOR FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: U.S. 60/357,383
PRIOR FILING DATE: 2002-02-15
                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/157,770
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: 60/163,939
PRIOR FILING DATE: 1999-11-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/11/058,926
CURRENT FILING DATE: 2005-02-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TITLE OF INVENTION: BOND CONTAINING PROTEINS IN FILE REFERENCE: HMV-052.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US/09/679,705
PRIOR FILING DATE: 2000-10-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: BECKWITH, JONATHAN APPLICANT: ASLUND, FREDRIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2002-(
NUMBER OF SEQ ID NOS: 18
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TITLE OF INVENTION: Method of Treating TRX Mediated Diseases
                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/166,044
                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: variant motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 4
TYPE: PRT
ORGANISM: homo sapiens
                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                        FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION:
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                                                                                         Similarity
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SEQ ID NOS: 24
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                                   CGPC 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RITZ, DANIEL
LIM, JACKIE EUN-AH
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                                                                         Conservative
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                                                                                         100.0%; Score 31; DB 7; 100.0%; Pred. No. 8.2e+04
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Pred. No. 8.2e+04;
                                                                         Mismatches
                                                                                       8.2e+04;
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Publication No. US20050260140A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: SEQ ID NO 2
                                         SOFTWARE: SEQ ID NO 3
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Best Local :
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                                                                                                        PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
PRIOR FILING DATE: 2005-02-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/462,082
PRIOR TILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
                                                                                         NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: White, Carl W. APPLICANT: del Val, Greg
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TYPE: PRI
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LOCATION: (1)..(6)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: synthetic peptide motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
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                                                                 PatentIn version 3.3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 31; DB 7; ilarity 100.0%; Pred. No. 8.2e+04; Conservative 0; Mismatches 0;
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Sequence 5372, Application US/10467657
Publication No. US20050260581A1
GENERAL INFORMATION:
APPLICANT: CHIRON SpA
APPLICANT: FONTANA MARIA RITA
APPLICANT: MASIGNANI Vega
APPLICANT: MASIGNANI Vega
APPLICANT: MONACI Eliabetta
APPLICANT: MONACI Eliabetta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: synthetic peptide motif US-11-090-916-3
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PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/669,373
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR FILING DATE: 2004-05-17
PRIOR FILING DATE: 2004-05-17
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
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Matches 4
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Publication No. US20060023808A1
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Best Local Similarity
Matches 4; Conserv
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LENGTH: 75
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APPLICANT: LEITE, JOHN F.
TITLE OF INVENTION: COMPOSITIONS, KITS, AI
TITLE OF INVENTION: SPECTROMETRY
TITLE REFERENCE: NVX-28 US
CURRENT EPPLICATION UNMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/621,686
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 9
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APPLICANT: POPE, ROBERT M.
APPLICANT: SADEGHI, MEHRNOOSH
CURRENT APPLICATION NUMBER: US/10/467,657
CURRENT FILING DATE: 2003-08-11
                                         TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Ver. 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: protein sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT ORGANISM: Artificial Sequence
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Pred. No.
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                                                                                                                                      US-11-074-176-286
                                                                                                                                                     RESULT 11
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CURRENT FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/572,290
PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/669,373
PRIOR FILING DATE: 2005-04-08
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR APPLICATION NUMBER: 60/621,686
PRIOR APPLICATION NUMBER: 60/621,686
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PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 9218
SOFTWARE: SeqWin99, version 1.04
SEQ ID NO 5372
LENGTH: 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 9
SOFTWARE: PATENTIN VEr. 3.3
SEQ ID NO 1
LENGTH: 91
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APPLICANT: HAJIVANDI, MAHBOD R.
                                                                                             Sequence 286, Appropriate Publication No.
                                                                             GENERAL INFORMATION:
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Best Local Similarity
Matches 4; Conserv
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Publication No. US20060023808A1
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APPLICANT: LEITE, JOHN F.
TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR CALIBRATION IN MASS
TITLE OF INVENTION: SPECTROMETRY
FILE REFERENCE: NVX-28 US
APPLICANT: Klaenhammer, Todd R. APPLICANT: Russell, William M. APPLICANT: Altermann, Eric APPLICANT: McAuliffe, Olivia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2004-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: protein sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
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11 Similarity 100.0%;
4; Conservative 0;
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                                                                                                                Application US/11074176
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100.0%; Pred. No.
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Pred. No.
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RESULT 13
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Sequence 7, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
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LENGTH: 103
TYPE: PRT
ORGANISM: Lactobacillus acidophilus
                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
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Best Local
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APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
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CURRENT APPLICATION NUMBER: US/11/074,176
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: 60/551,161
PRIOR FILING DATE: 2004-03-08
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NUMBER OF SEQ ID NOS: 15
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PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10 PRIOR APPLICATION NUMBER: 60/462,082
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CURRENT FILING DATE: 2005-03-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 4.0
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TITLE OF INVENTION: Nucleic Acid Sequences Encoding
TITLE OF INVENTION: Stress-Related Proteins and Uses Therefore
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/650,865
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                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORĜANISM: Listeria monocytogenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
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                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 31; DB 7; Length 103; 100.0%; Pred. No. 32;
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Best Local Similarity
Thes 4; Conserve
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PRIOR FILING DATE: 2003-04-11
PRIOR PELICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.3 SEQ ID NO 5
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                                                                           Matches
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Best Local (
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TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
CURRENT FILING DATE: 2005-03-24
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CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILLING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILLING DATE: 2002-09-10
PRIOR FILLING DATE: 2002-09-10
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PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
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                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
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TYPE: PRT
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TYPE: PRT
                                                                                           Local Similarity
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nilarity 100.0%;
Conservative 0
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Pred. No.
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US-10-821-234-1371 US-10-821-234-1371 ; Sequence 1371, Application US/10821234 ; Publication No. US20050255114A1

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                                                                                             RESULT 17
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US-10-821-234-1371
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US-11-090-916-8
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CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
SOFTWARE: pt SEQ genes Version 1.0
SEQ ID NO 1371
Sequence 9, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 8, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.3
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PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
PRIOR FILING DATE: 2004-03-24
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PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
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TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
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TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
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CURRENT APPLICATION NUMBER: US/11/090,916

CURRENT FILING DATE: 2005-03-24

PRIOR APPLICATION NUMBER: 60/440,960

PRIOR FILING DATE: 2002-09-10

PRIOR APPLICATION NUMBER: 60/462,082

PRIOR FILING DATE: 2003-04-11

PRIOR APPLICATION NUMBER: 10/660,118

PRIOR PILING DATE: 2003-09-10

PRIOR PRIOR FILING DATE: 2003-09-10

PRIOR APPLICATION NUMBER: 60/556,516

PRIOR APPLICATION NUMBER: 60/556,516

PRIOR APPLICATION NUMBER: 60/650,865

PRIOR APPLICATION NUMBER: 60/650,865

PRIOR FILING DATE: 2005-02-07
                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.3 SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                APPLICANT: del Val, Greg
APPLICANT: del Val, Greg
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR FILING DATE: 2002-09-11
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2003-09-10
PRIOR PILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
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SEQ ID NO 9
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PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
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                                                                                                                                                                         TYPE: PRT ORGANISM: Rattus norvegicus
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; ORGANISM: Homo sapiens
US-11-090-916-12
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US-11-090-916-11
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SEQ ID NO 12
LENGTH: 105
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                                                        Query Match 100.0%;
Best Local Similarity 100.0%;
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PRIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
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PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
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TITLE OF INVENTION: MucOlytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR RILING DATE: 2002-09-10
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PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
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PRIOR PILLING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
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APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
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NUMBER OF SEQ ID NOS: 15
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                                    Conservative
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Pred. No.
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US-11-144-301A-10
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US-11-144-301A-9
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US-11-106-796-13
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LENGTH: 105
TYPE: PRT
Sequence 10, Application US/11144301A Publication No. US20050288227A1 GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 9
LENGTH: 105
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Best Local Similarity
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CURRENT FILING DATE: 2005-04-15
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/577,089
PRIOR FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 60/357,383
PRIOR FILING DATE: 2002-02-15
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APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF PERTILITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Johanna, Ungerstedt
TITLE OF INVENTION: USE OF THIOREDOXIN MEASUREMENTS FOR DIAGNOSTICS AND TREATMENTS
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APPLICANT: Marks, Paul A.

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US-11-032-773-943
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NUMBER OF SEQ ID NOS: 958
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 943
                                                                         Sequence 19, Application US/10689742 Publication No. US20050250180A1 GENERAL INFORMATION:
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Publication No. US20
GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity
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TITLE OF INVENTION: DESIGN OF THERAPEUTICS
FILE REFERENCE: 17102-013001 / 1762
CURRENT APPLICATION NUMBER: US/11/032,773
CURRENT FILING DATE: 2005-01-11
PRIOR APPLICATION NUMBER: 60/536,184
PRIOR FILING DATE: 2004-01-12
PRIOR APPLICATION NUMBER: 60/557,591
PRIOR APPLICATION NUMBER: 60/557,591
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PRIOR FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 60/357,383
PRIOR FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 10
APPLICANT: Jacobs, Kenneth
APPLICANT: McCoy, John M
APPLICANT: LaVallie, Edward R
APPLICANT: Racie, Lisa A
                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Mus musculus
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank BAA04881
DATABASE ENTRY DATE: 2002-12-25
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TITLE OF INVENTION: USE OF THIOREDOXIN MEASUREMENTS FOR DIAGNOSTICS AND TREATMENTS
FILE REFERENCE: 28517-501 UTIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo sapiens
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100.0%; Pred. No. 32;
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APPLICANT: Kagur, Vivek and Gebhart, Connie J.
TITLE OF INVENTION: NUCLEIC ACID AND POLYPEPTIDE SEQUENCES
TITLE OF INVENTION: FROM LAWSONIA INTRACELLULARIS AND METHODS
FILE REFERENCE: 09531-128001
CURRENT APPLICATION NUMBER: US/11/098,686
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US03/31318
PRIOR APPLICATION NUMBER: PCT/US03/31318
PRIOR FILING DATE: 2003-10-01
PRIOR FILING DATE: 2003-10-01
PRIOR FILING DATE: 2003-10-04
NUMBER OF SEQ ID NOS: 11433
SOPTWARE: FRASESEQ for Windows Version 4.0
SEQ ID NO 10612
LENGTH: 107
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US-11-098-686-10612
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Sequence 4, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
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Best Local S
Matches 4
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Publication No. US20060024696A1
GENERAL INFORMATION:
APPLICANT: Kapur, Vivek and Gebhart, Co
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CURRENT FILING DATE: 2003-10-22
PRIOR APPLICATION NUMBER: 09/746,783
PRIOR FILING DATE: 2000-12-21
NUMBER OF SEQ ID NOS: 231
SOFTWARE: Patentin version 3.2
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APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
FILE REFERENCE: 00766.000091.10
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TYPE: PRT
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                                                                                                                      ; TYPE: PRT ; ORGANISM: Homo sapiens US-11-072-512-3199
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SEQ ID NO 4
LENGTH: 109
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SEQ ID NO 3199
LENGTH: 113
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Best Local
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Best Local Similarity
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TITLE OF INVENTION: Novel full length cDNA
FILE REFERENCE: 084335-0191
CURRENT APPLICATION UMBER: US/11/072,512
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: US 60/350,978
PRIOR FILING DATE: 2002-01-25
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NUMBER OF SEQ ID NOS: 4096
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PRIOR FILING DATE: 2005-02-07
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PRIOR APPLICATION NUMBER: JP 2001-379298
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PRIOR FILING DATE: 2003-04-11
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FILING DATE: 2004-03-24
37 CGPC 40
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OTSUKI, TETSUJI
WAKAMATSU, AI
SATO, HIROYUKI
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YOSHIKAWA, TSUTOMU
OTSUKA, MOTOYUKI
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NAGAI, KEIICHI
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TAMECHIKA, ICHIRO
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YAMAMOTO, JUN-ICHI
                                                               Conservative
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100.0%; Pred. No. 34;
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RESULT 29
US-11-195-459-12
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; TYPE: PRT
; ORGANISMF Hordeum vulgare
US-11-195-459-16
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; TYPE: PRT
; ORGANISM: Zea mays
US-11-195-459-12
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Publication No. US20050278803A1
                                                                                                                                                                                                              SEQ ID NO 16
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Best Local Similarity
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Best Local Similarity
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APPLICANT: Everard, John
APPLICANT: Allen, Stephen
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PROFILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
PRIOR FILING DATE: 2001-12-03
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PRIOR FILING DATE: 2001-12-03
PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
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PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn version 3.0
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CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
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APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PROTITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PROTITURE OF INVENTION:
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    2 CGPC 5
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Meeley, Robert
Hantke, Sabine
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Hantke, Sabine
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                                         100.0%; Score 31; DI
100.0%; Pred. No. 36
Live 0; Mismatches
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Pred. No. 36;
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                                                                                    DB 7; Length 122;
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SEQ ID NO 10
LENGTH: 123
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Query Match
Best Local Similarity
Matches 4; Conserv
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SEQ ID NO 19
LENGTH: 122
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                                                                                                                                                                                                                                                         APPLICANT: Sewalt, Vincent
APPLICANT: Hastings, Craig
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PR
TITLE OF INVENTION NUMBER: US/11/195,459
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2001-12-03
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR APPLICATION NUMBER: US/10/005,429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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Best Local (
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE PILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
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PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
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FEATURE:
RAME/KEY: misc_feature
OTHER INFORMATION: "n" at position 9, 493, 537, 548,
OTHER INFORMATION: or t
                                                                                                       TYPE: PRT
ORGANISM: Zea mays
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No. US20050278803A1
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    100.0%; Score 31; DB 7; Length 123; ilarity 100.0%; Pred. No. 36; Conservative 0; Mismatches 0; Indels
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Best Local
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RESULT 35
US-11-106-796-11
Squence 11, Application US/11106796
Publication No. US20050282191A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: SUTOVSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO
APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Homo sapiens US-10-995-561-703
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US-10-995-561-703
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PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 10
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 10, Application US/11106796
Publication No. US20050282191A1
GENERAL INFORMATION:
APPLICANT: SUTOVSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 703
LENGTH. 126
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
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TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE,
TITLE OF INVENTION: DETECTION AND USES THEREOF
FILE REFERENCE: CL001559
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: UVMO:033US
CURRENT APPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2005-04-15
                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 127
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b. US20050272054A1
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100.0%; Pred. No. 37
tive 0; Mismatches
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US-11-106-796-12
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; TYPE: PRT
; ORGANISM: Mus musculus
US-11-106-796-11
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SEQ ID NO 12
LENGTH: 127
SOFTWARE: SeqWin99,
SEQ ID NO 4354
LENGTH: 129
TYPE: PRT
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Best Local Similarity
Matches 4; Conser
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Best Local Similarity
                                                               CURRENT APPLICATION NUMBER: US/10/467,657
CURRENT FILING DATE: 2003-08-11
PRIOR APPLICATION NUMBER: GB-0103424.8
PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 9218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: SUTOVSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO.

RITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
FILE REFERENCE: UVMO:033US
CURRENT APPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2005-04-15
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR PILICATION NUMBER: 60/562,526
RIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
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CURRENT FILING DATE: 2005-04-15
PRIOR APPLICATION NUMBER: 60/562,526
'PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                            APPLICANT: PIZZA Mariagrazia
APPLICANT: MASIGNANI Vega
APPLICANT: MONACI Eliaabetta
TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS
                                                                                                                                                                                                                                                 APPLICANT: CHIRON SpA
APPLICANT: FONTANA Maria Rita
                                                                                                                                                                 FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: RAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: UVMO:033US
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Pred. No. 37;
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Pred. No. 37;
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US-11-090-916-13
Sequence 13, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 205-03-24
CURRENT FILING DATE: 205-03-24
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Best Local Similarity
---hes 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: MASUHO, YASUHIKO
TITLE OF INVENTION: NOVEL full length cDNA
PILE REFERENCE: 084335-0191
CURRENT APPLICATION NUMBER: US/11/072,512
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: US 60/350,978
PRIOR FILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: JP 2001-379298
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 4096
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US-11-072-512-2145
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Matches
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SQ ID NO 2145
LENGTH: 129
TYPE: PRT
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PRIOR APPLICATION NUMBER: 60/409,960
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                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                     Local Similarity
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NAGAI, KEIICHI
IRIE, RYOTARO
TAMECHIKA, ICHIRO
SEKI, NAOHIKO
YOSHIKAWA, TSUTOMU
OTSUKA, MOTOYUKI
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OTSUKI, TETSUJI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAGAHARI, KENJI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WAKAMATSU, AI
SATO, HIROYUKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               YAMAMOTO, JUN-ICHI
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Conservative 0; Mismatches
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Pred. No.
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37;
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RESULT 40
US-11-195-459-8
; Sequence 8, Application US/11195459
; Publication No. US20050278803A1
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                                                                                                                           RESULT 41
US-09-978-360A-641
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US-11-195-459-8
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Best Local S
Matches 4
                                         Sequence 641, Application US/09978360A
Publication No. US20060009633A9
GENERAL INFORMATION:
APPLICANT: Edwards, Jean-Baptiste Dumas Milne
APPLICANT: Duclert, Aymeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Sewalt,
APPLICANT: Hasting
APPLICANT: Meeley,
APPLICANT: Hantke,
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                                                                                                                                                                                                                                                                              Query Match
Best Local
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SEQ ID NO 8
                                                                                                                                                                                                                                                               Matches
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APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF
FILE REFERENCE: 5718-119 (035718/241421)
CURRENT FILLORION NUMBER: US/11/195,459
CURRENT FILLING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR PILLING DATE: 2001-12-03
PRIOR FILLING DATE: 2001-12-03
PRIOR FILLING DATE: 2001-12-03
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PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
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            APPLICANT:
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PRIOR APPLICATION 1
APPLICANT:
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TYPE: PRT
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APPLICATION NUMBER: 60/650,865
FILING DATE: 2005-02-07
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4; Conserva
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Meeley, Robert
Hantke, Sabine
            Duclert, Aymeric
Bougueleret, Lydie
Jobert, Severin
                                                                                                                                                                                                                                                          100.0%; ilarity 100.0%; Conservative 0
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Pred. No.
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38;
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US-11-090-916-14
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; LOCATION: -41..-1
US-09-978-360A-641
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CURRENT FILING DATE: 2001-10-15
PRIOR APPLICATION NUMBER: US 60/066,677
PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: US 60/069,957
PRIOR APPLICATION NUMBER: US 60/074,121
PRIOR APPLICATION NUMBER: US 60/074,121
PRIOR FILING DATE: 1998-02-09
PRIOR APPLICATION NUMBER: US 60/081,563
PRIOR APPLICATION NUMBER: US 60/096,116
PRIOR FILING DATE: 1998-04-13
PRIOR APPLICATION NUMBER: US 60/096,116
PRIOR FILING DATE: 1998-08-10
PRIOR APPLICATION NUMBER: US 60/096,116
PRIOR APPLICATION NUMBER: US 60/099,273
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Remaining Prior Application data removed - See File Wrapper or PALM.

NUMBER OF SEQ ID NOS: 810

SOFTWARE: Patent.pm

SEQ ID NO 641

LENGTH: 143

TYPEF The Patent of t
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Matches
SEQ ID NO 14
LENGTH: 167
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                                                                                                                                                                          TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof FILE REFERENCE: 2879-98-1 CURRENT APPLICATION NUMBER: US/11/090,916 CURRENT FILING DATE: 2005-03-24 PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10 PRIOR FILING DATE: 2003-04-11 PRIOR APPLICATION NUMBER: 60/462,082 PRIOR FILING DATE: 2003-04-11 PRIOR APPLICATION NUMBER: 60/60,118 PRIOR APPLICATION NUMBER: 60/550,516 PRIOR FILING DATE: 2003-04-10 PRIOR FILING DATE: 2004-03-24 PRIOR FILING DATE: 2004-03-24 PRIOR PRIOR APPLICATION NUMBER: 60/550,516 PRIOR FILING DATE: 2004-03-24 PRIOR PILING DATE: 2004-03-24 PRIOR FILING DATE: 2004-03-24 PRIOR FILING DATE: 2004-03-24 PRIOR FILING DATE: 2005-02-07 NUMBER: 60/550,865 PRIOR FILING DATE: 2005-02-07 NUMBER: 60/550,865 PRIOR FILING DATE: 2005-03-10 PRIOR FILING DATE: 2005-03-04 PRIOR P
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APPLICANT: del Val, (
APPLICANT: Lee, Rees
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PRIOR FILING DATE: 1998-11-13
PRIOR APPLICATION NUMBER: US 09/215,435
PRIOR FILING DATE: 1998-12-17
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                                                                                                                                   SOFTWARE: PatentIn version 3.3
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FILING DATE: 1998-12-17
APPLICATION NUMBER: US 09/247,155
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ORGANISM: Zea mays

TYPE: PRT

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US-11-090-916-15
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PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR ETITIOR DATE: 2004-09-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.3 SEQ ID NO 15 LENGTH: 172
                                                                                                                                                                                                                                                                                                                           Sequence 3, Application US/11131744
Publication No. US20060023808A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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APPLICANT:
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Best Local Similarity
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                              PRIOR APPLICATION NUMBER: 60/572,290
PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/669,373
PRIOR FILING DATE: 2005-04-08
PRIOR APPLICATION NUMBER: 60/572,215
                                                                                                                   APPLICANT: LIANG, XIQUAN
APPLICANT: LEITE, JOHN F.
TITLE OF INVENTION: COMPOSITIONS, KITS, AN
TITLE OF INVENTION: SPECTROMETRY
FILE REFERENCE: NVX-28 US
CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: White, Carl W. APPLICANT: del Val, Greg APPLICANT: Lee, Rees
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
                                                                                                                                                                                                                                                                        APPLICANT: HAJIVANDI, MAHBOD R. APPLICANT: POPE, ROBERT M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/650,865 PRIOR FILING DATE: 2005-02-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Mucolytic and Anti-Blastase Compounds and Methods of Use Thereof FILE REFERENCE: 2879-98-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Oryza sativa
FILING DATE: 2004-05-17
APPLICATION NUMBER: 60/621,685
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 2004-03-24
                                                                                                                                                                                                                                                                                                                                                                                                                                         95 CGPC 98
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Pred. No.
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RESULT 46

US-10-995-561-643

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(US-10-995-561-643)

(US-10-995-561-643)

(US-10-995-561)

(US-10-995-961)

(US-10-995-961)
; ORGANISM: Homo sapiens US-10-995-561-643
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                                                 NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 643
LENGTH: 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Plesch, Gunnar
APPLICANT: Blau, Astrid
APPLICANT: Laeschner, Klaus
APPLICANT: Caeschner, Klaus
APPLICANT: Caeschner, Klaus
APPLICANT: Color Michael (1970)
TITLE OF INVENTION: Identification of Herbicidally Active Substances
FILE REFERENCE: 2000 857
CURRENT FILING DATE: 2003-08-14
PRIOR APPLICATION NUMBER: PCT/EP02/01466
PRIOR FILING DATE: 2002-02-13
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Best Local Similarity
Matches 4; Conserv
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SEQ ID NO 3
LENGTH: 177
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Best Local Similarity 100.0%; Pred. No.
Matches 4; Conservative 0; Mismatch
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TYPE: PRT
ORGANISM: Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/621,686
PRIOR FILING DATE: 2004-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
                                           TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: protein sequence
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GENETIC POLYMORPHISMS ASSOCIATED WITH
CARDIOVASCULAR DISORDERS AND DRUG RES
DETECTION AND USES THEREOF
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Pred. No.
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CURRENT APPLICATION NUMBER: US/11/064,774A
CURRENT FILING DATE: 2005-02-24
PRIOR APPLICATION NUMBER: 09/795,006
PRIOR FILING DATE: 2001-02-26
PRIOR PLICATION NUMBER: US 60/205,331
PRIOR FILING DATE: 2000-05-18
PRIOR APPLICATION NUMBER: US 60/185,205
PRIOR APPLICATION NUMBER: US 60/185,205
PRIOR APPLICATION NUMBER: US 60/185,205
PRIOR FILING DATE: 2000-02-25
NUMBER OF SEQ ID NOS: 1212
SOFTWARE: PATENTIN Ver. 3.2
SEQ ID NO 2
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US-10-995-561-642
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                                                                                                                                            ; TYPE: PRT; ORGANISM: Homo sapiens US-11-064-774A-2
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Best Local S
Matches 4
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GENERAL INFORMATION:
APPLICAMT: CARGILL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
TITLE OF INVENTION: DETECTION AND USES THEREOF
TITLE REFERENCE: CL001559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT APPLICATION UNMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOPTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 642
**PROPRIET 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/11064774A Publication No. US20050267024A1 GENERAL INFORMATION:
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Best Local Similarity
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                                                                        Matches
                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: MATERIALS AND METHODS INVOLVING HYBRID VASCULAR ENDOTHELIAL GROWTITLE OF INVENTION: FACTOR DNAS AND PROTEINS
FILE REFERENCE: 28967/35977B2
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                                                                                                                                                                                                   LENGTH: 191
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 143 CGPC 146
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                                                                        Conservative
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100.0%; Pred. No.
                                                                                         100.0%;
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                                                                                         Score 31; DB
Pred. No. 47;
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47;
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                                                                                                        Length 191;
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; SEQ ID NO 2
, LENGTH: 191
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-075-400-2
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US-11-226-005-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 4, Application US/11226005
Publication No. US20060014252Al
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: ERIKKSON, et al.
TITLE OF INVENTION: MULTIVALENT ANTIBODY MATERIALS AND METHODS FORVEGF/PDGF FAMILY OF
TITLE OF INVENTION: GROWTH FACTORS
FILE REFERENCE: 28967/39820B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 32
           TELEX: 756822
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                            COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.
SOFTWARE: Microsoft Word, Version 5.
APPLICATION NUMBER: US/11/226,005
FILING DATE: 14-Sep-2005
FILING DATE: 14-Sep-2005
FILING DATE: 05-NOV-196
APPLICATION UNMBER: US/08/743,868
FILING DATE: 05-NOV-196
APPLICATION UNMBER: 08/54,374
FILING DATE: 08-NOV-195
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
les 4; Conserv
                                                                                                        NAME: Anderson, Kathryn A.
REGISTRATION NUMBER: 32,172
REFERENCE/DOCKET NUMBER: 2837
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Lyman, Stewart D. et al.
TITLE OF INVENTION: flk-1 Binding Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             143 CGPC 146
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Immunex Corporation STREET: 51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Seattle
                                                                   TELEPHONE: (200, 233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: US
LENGTH: 191 amino acids
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; ORGANISM: Bovine US-11-145-494-2
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-129-076-9
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US-11-129-076-9
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Best Local Similarity luv.
"""ches 4; Conservative
                                 SEQ ID NO 2
LENGTH: 191
TYPE: PRT
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APPLICANT: Edelberg, Jay A.
APPLICANT: Xaymarden, Munirva
                                                                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/11145494
Publication No. US20060024785A1
GENERAL INFORMATION:
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Publication No. US20060019891A1
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APPLICANT: Cormell Research Foundation, I
TITLE OF INVENTION: Protection of Cardiac
FILE REFERENCE: 1676.009US1
CURRENT APPLICATION NUMBER: US/11/129,076
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 18
                                                                                                                                                                                                                              FILE REFERENCE: A-62326-2
CURRENT APPLICATION NUMBER: US/11/145,494
CURRENT FILING DATE: 2005-06-03
                                                                                                                                                                                                                                                                          APPLICANT: Keyt, Bruce A.
APPLICANT: Nguyen, Francis H.
APPLICANT: Ferrara, Napoleone
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING VARIANTS OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: CELL GROWTH FACTOR
                                                                                      NUMBER OF SEQ ID NOS: 42
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                     PRIOR FILING DATE: 1995-12-05
PRIOR APPLICATION NUMBER: 60/002,827
PRIOR FILING DATE: 1995-08-25
                                                                                                                                                                       PRIOR APPLICATION NUMBER: US/09/346,069
PRIOR FILING DATE: 1999-07-01
PRIOR APPLICATION NUMBER: 08/567,200
PRIOR APPLICATION NUMBER: 08/567,200
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CLONE: Hu VEGF
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
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PRIOR APPLICATION NUMBER: US 60/416,395
PRIOR FILING DATE: 2002-10-04
NUMBER OF SEQ ID NOS: 11433
SOPTWARE: FRACTSC for Windows Version 4.0
SEQ ID NO 11017
LENGTH: 197
TYPE: PRT
ORGANISM: Lawsonia intracellularis
US-11-098-686-11017
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US-11-098-686-11017
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US-11-075-047A-81
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Publication No. US20060024696A1

GENERAL INFORMATION:

APPLICANT: Kapur, Vivek and Gebhart, Connie J.

APPLICANT: NUCLBIC ACID AND POLYPEPTIDE SEQUENCES

TITLE OF INVENTION: FROM LAWSONIA INTRACELLULARIS AND METHODS OF USING

FILE REFERENCE: 09531-128001

CURRENT APPLICATION NUMBER: US/11/098,686

CURRENT FILING DATE: 2005-04-04

PRIOR APPLICATION NUMBER: PCT/US03/31318

PRIOR PILING DATE: 2003-10-01
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 81
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Publication No. US20060030000A1
GENERAL INFORMATION:
APPLICANT: ALITALO, et al.
TITLE OF INVENTION: GROWTH FACTOR BINDING CONSTRUCTS MATERIALS AND METHODS
FILE REFERENCE: 28967/39700A
                                                                                          Query Match
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Best Local Similarity
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                                                                              Local Similarity
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                             2 CGPC 5
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                                                            4; Conservative
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                                                                             Score 31;
Pred. No.
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RESULT 55 US-10-467-657-8494 ; Sequence 8494, Application US/10467657

Publication No. US20050260581A1

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GENERAL INFORMATION:

APPLICANT: CARGILL, Michele et al.

TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH

TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF

TITLE OF INVENTION: DETECTION AND USES THEREOF

FILE REFERENCE: CL001559

CURRENT APPLICATION NUMBER: US/10/995,561

CURRENT FILING DATE: 2004-11-24

NUMBER OF SEQ ID NOS: 85702

SOFTWARE: FRSESEQ for Windows Version 4.0

SEQ ID NO 640

LENGTH: 209

TYPE: PRT

ORGANISM: Homo sapiens
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Best Local Similarity
"---hes 4; Conserve
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US-10-995-561-645
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; ORGANISM: Neisseria gonorrhoeae
US-10-467-657-8494
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                                                                                                   Sequence 645, Application US/10995561

Publication No. US20050272054A1

GENERAL INFORMATION:

APPLICANT: CARGILL, Michele et al.

TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH

TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS

TITLE OF INVENTION: DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 9218
SOFTWARE: SeqWin99, version 1.04
SEQ ID NO 8494
LENGTH: 209
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 640, Appropriate Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: CHIRON SPA
APPLICANT: FONTANA MARIA RITA
APPLICANT: FIZZA MARIAGRAZIA
APPLICANT: MASIGNANI Vega
APPLICANT: MONACI Elisabetta
TITLE OF INVENTION: GONOCOCCCAL PROTEINS AND NUCLEIC ACIDS
FILE REFERENCE: CL001559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/467,657
CURRENT FILING DATE: 2003-08-11
PRIOR APPLICATION NUMBER: GB-0103424.8
PRIOR FILING DATE: 2001-02-12
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nes 4; Conserv
                                                                                                                                                                                                                                                                                                                                                 161 CGPC 164
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10. US20050272054A1
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100.0%; Pred. No. 50
Live 0; Mismatches
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                                                                       US-11-149-462-3
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Publication OS. US20050272054A1
GENERAL INFORMATION:
APPLICANT: CARGILL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CARGILLAR DISORDERS AND DRUG RESPONSE, METHODS:
TITLE OF INVENTION: DETECTION AND USES THEREOF
FILE REFERENCE: CL001559
CURRENT APPLICATION UNBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOCTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 646
LENGTH: 215
TYPE: DET
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Best Local Similarity 100.
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LENGTH: 215
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LENGTH: 215
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Best Local Similarity
            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                      APPLICANT: Shea, Lonnie D.
APPLICANT: Shea, Lonnie D.
APPLICANT: Whittlesey, Kevin
APPLICANT: Whittlesey, Kevin
APPLICANT: Yang, Yang
APPLICANT: Rives, Christopher
APPLICANT: Rovedo, Mark
APPLICANT: Iskandar, Bermans
TITLE OF INVENTION: Biodegradable Scaffolds and Uses Thereof
FILE REFERENCE: 1720-1-011N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
   Matches
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PRIOR FILING DATE: 2004-06-10
NUMBER OF SEQ ID NOS: 12
SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2005-06-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Homo sapiens
                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
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 Conservative
100.0%; Score 31;
100.0%; Pred. No.
tive 0; Mismatch
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US-11-106-399-8
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; ORGANISM: Homo sapiens US-11-072-512-3343
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CURRENT APPLICATION NUMBER: US/11/106,399
CURRENT FILING DATE: 2005-04-14
NUMBER OF SEQ ID NOS: 10
SOPTWARE: PATENTIN VET. 2.1
SEQ ID NO 8
LENGTH: 216
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Publication No. US20060002892A1
GENERAL INFORMATION:
APPLICANT: MATHEW, PORUNELLOOR A.
APPLICANT: BOLES, KENT S.
TITLE OF INVENTION: LLT USES THEREOF IN IMMUNE SYSTEM MODULATION
                          SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3343
LENGTH: 224
TYPE: PRT
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Best Local Similarity
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Publication No. US20060029945A1
                                                                                                                                                                               APPLICANT: NAGAHARI, KENJI
APPLICANT: WASUHO, YASUHIKO
TITLE OF INVENTION: Novel full length cDNA
FILE REFERENCE: 084335-0191
CURRENT APPLICATION NUMBER: US/11/072,512
CURRENT FILING DATE: 2005-03-07
                                                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
                                                                                               PRIOR APPLICATION NUMBER: JP 2001-379298
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 4096
                                                                                                                                               CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: US 60/350,978
PRIOR FILING DATE: 2002-01-25
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ORGANISM: Homo sapiens
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SATO, HIROYUKI
ISHII, SHIZUKO
YAMAMOTO, JUN-ICHI
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NAGAI, KEIICHI
IRIE, RYOTARO
TAMECHIKA, ICHIRO
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YOSHIKAWA, TSUTOMU
OTSUKA, MOTOYUKI
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OTSUKI, TETSUJI
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RESULT 64 US-11-153-880-7

Sequence 7, Application US/11153880 Publication No. US20050256050A1 GENERAL INFORMATION:

APPLICANT:

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APPLICANT: CARGILL, Michele et al.
APPLICANT: CARGILL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATE
TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND I
TITLE OF INVENTION: DETECTION AND USES THEREOF
FILE REFERENCE: CL001559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 641
LENGTH: 232
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CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 644
LENGTH: 232
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Best Local Similarity 100.0%;
Matches 4; Conservative 0
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Best Local
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Best Local Similarity
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TITLE OF INVENTION: CARI
TITLE OF INVENTION: DETI
FILE REFERENCE: CL001559
                                                                                                                                                                          LENGTH: 23
TYPE: PRT
                                                                                                                                                          ORGANISM: Homo sapiens
184 CGPC 187
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                                                                  100.0%; Score 31; DB nilarity 100.0%; Pred. No. 53; Conservative 0; Mismatches
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GENETIC POLYMORPHISMS ASSOCIATED WITH
CARDIOVASCULAR DISORDERS AND DRUG RE
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GENETIC POLYMORPHISMS ASSOCIATED WITH
CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS
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Pred. No. 52;
0; Mismatches
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                                                                                                      DB 6; Length 232;
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                                                                                                                                                                                                                    Publication No. US20050267024A1
GENERAL INFORMATION:
APPLICANT: Alitalo et al.
                                                                                                                                                                                                                                                                       Sequence 147, Application US/11064774A
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             CURRENT APPLICATION NUMBER: US/11/064,774A CURRENT FILING DATE: 2005-02-24 PRIOR APPLICATION NUMBER: 09/795,006 PRIOR FILING DATE: 2001-02-26 PRIOR APPLICATION NUMBER: US 60/205,331 PRIOR FILING DATE: 2000-05-18 PRIOR APPLICATION NUMBER: US 60/185,205 PRIOR APPLICATION NUMBER: US 60/185,205 PRIOR FILING DATE: 2000-02-25
                                                                                                                                                              TITLE OF INVENTION: MATERIALS AND METHODS INVOLVING HYBRID VASCULAR ENDOTHELIAL GROWTITLE OF INVENTION: FACTOR DNAS AND PROTEINS
FILE REFERENCE: 28967/35977B2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (202)371-2540 [INFORMATION FOR SEQ ID NO: 7: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: VASCULAR ENDOTHELIAL GROWTH FACTOR 2
NUMBER OF SEQUENCES: 15
CORRESCE: STERNE VECC.
STEREFT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA RELEASE #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08
FILING DATE: 06-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: MARKOWICZ, KAREN R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
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REFERENCE/DOCKET NUMBER: 1488.1000004
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)371-2600
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APPLICATION NUMBER:
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APPLICATION NUMBER:
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SEQ ID NOS: 1212
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        232 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 7;
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; TYPE: PRT ; ORGANISM: Homo sapiens US-11-064-774A-147
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US-11-233-119-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-11-211-724-5
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SEQ ID NO 147
LENGTH: 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 5
LENGTH: 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 5, Application US/11211724
Publication No. US20050287143A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                    SOFTWARE: PatentIn version 3.0
SEQ ID NO 7
LENGTH: 232
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7, Application US/11233119 Publication No. US20060025331A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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CURRENT FILING DATE: 2005-09-23
PRIOR APPLICATION NUMBER: 09/107,997
PRIOR FILING DATE: 1998-06-30
PRIOR APPLICATION NUMBER: 09/042,105
PRIOR APPLICATION NUMBER: 09/042,105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US/08/465,968
PRIOR FILING DATE: 1995-06-06
PRIOR APPLICATION NUMBER: 08/207,550
PRIOR FILING DATE: 1994-03-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.0
                                                                                                                    PRIOR APPLICATION NUMBER: 08/465,968
PRIOR TILING DATE: 1995-06-06
PRIOR APPLICATION NUMBER: 08/207,550
PRIOR FILING DATE: 1994-03-08
                                                                                                                                                                                                   PRIOR FILING DATE: 1998-03-13
PRIOR APPLICATION NUMBER: 09/999,811
PRIOR FILING DATE: 1997-12-24
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: HU, JING-SHAN et al,
TITLE OF INVENTION: VASCULAR ENDOTHELIAL GROWTH FACTOR 2
FILE REFERENCE: PF112P4D1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: PF112P1
CURRENT APPLICATION NUMBER: US/11/211,724
CURRENT FILING DATE: 2005-08-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Vascular Endothelial Growth Factor 2
                                                                                                     NUMBER OF SEQ ID NOS: 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: homo sapiens
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Pred. No.
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US-11-233-119-7

Query Match Best Local Similarity

100.0%; Score 31; DB 7; Length 232; 100.0%; Pred. No. 53;

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; NUMBER OF SEQ ID NOS: 128
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 91
; LENGTH: 232
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-075-047A-91
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US-11-131-744-6
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Best Local Similarity lov.
"Atches 4; Conservative
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Matches
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                                                                           SOFTWARE: PatentIn Ver. 3.3
SEQ ID NO 6
LENGTH: 273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6, Application US/11131744 Publication No. US20060023808A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: ALITALO, et al.
TITLE OF INVENTION: GROWTH FACTOR BINDING CONSTRUCTS MATERIALS AND METHODS
FILE REFERENCE: 29867/39700A
CURRENT APPLICATION NUMBER: US/11/075,047A
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: US 60/550,907
PRIOR PILIAGE DATE: 2004-03-07
                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/572,290
PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/669,373
PRIOR FILING DATE: 2005-04-08
PRIOR FILING DATE: 2005-04-08
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: POPE, ROBERT M.
APPLICANT: SADECHI, MEHRNOOSH
APPLICANT: LIANG, XIQUAN
APPLICANT: LEITE, JOHN F.
TITLE OF INVENTION: SPECTROMETRY
TITLE OF INVENTION: SPECTROMETRY
                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: HAJIVANDI, MAHBOD R. APPLICANT: POPE, ROBERT M.
                                                                                                                                         NUMBER OF SEQ ID NOS:
                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/621,685
PRIOR FILING DATE: 2004-10-26
                                                                                                                                                            PRIOR FILING DATE: 2004-10-26
                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/572,215
PRIOR FILING DATE: 2004-05-17
                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: NVX-28 US
                    ORGANISM: Artificial Sequence FEATURE:
                                                          TYPE: PRT
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
                                                                                                                                                                               APPLICATION NUMBER: 60/621,686
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Search completed: February 23, 2006, 00:48:08
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                                                                                                                                                   US-11-186-284-85
                                                                                                                                                                                  SEQ ID NO 85
LENGTH: 282
TYPE: PRT
                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US/10/301,822
PRIOR FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US 60/339,971
PRIOR FILING DATE: 2001-12-10
PRIOR APPLICATION NUMBER: US 60/361,978
PRIOR APPLICATION NUMBER: US 60/361,988
PRIOR APPLICATION NUMBER: US 60/381,988
PRIOR APPLICATION NUMBER: US 60/381,988
PRIOR FILING DATE: 2002-05-20
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Best Local Similarity 100
Matches 4; Conservative
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Publication No.
                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: MPMO1-029P2RNM
CURRENT APPLICATION NUMBER: US/11/186,284
CURRENT FILING DATE: 2005-07-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Thibodeau, Stephen N.
APPLICANT: Burgart, Lawrence J.
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND
TITLE OF INVENTION: METHODS FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND
TITLE OF INVENTION: THERAPY OF COLON CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Millennium Pharmaceuticals, Inc. APPLICANT: Berger, Allison
                                                                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                   ORGANISM: Homo Sapiens
                                                                                           Local Similarity
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2 CGPC 35
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Kamatkar, Shubhangi
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Monahan, John E.
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No. US20050266493A1
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tive 0; Mismatches
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Pred. No.
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Abb07837 Aau65827	bg71301	dc53945 ae05156	ab93204 am94022	ab95309	dp55389	eb14143	dz03397 dz52354	du50414 dw49693	dt03346	di96410	dm28376 di95858	dm42514	dm17458 d107292	dm27790	dk82892	dj63637 dj77532	di18356	dg09306 di14761	di15429	dh25681	dg62656	dh43547	dg59791	dh28606 dq54751	dg48760 dg61519	dh12673	dg81168 dg56407	dg49384	dg52471	dg50008 dg51880	dh12049	dg81720	.dg71533 .dq50632	dg53647	dg58063	dg58615	.dg55855	dg57511	.dg62071 .dg82272	dg60967	Adg24959 Adg07256 Adg07808
Amino		Huma	Huma	Human Human	Human	Cance	Human Human	PRO101 PRO101	Human	Novel	Human Human	Human	Human	Human	Human	Novel Human	Novel	Novel	Nove1	Human	Human	Human	Novel	Human Novel	Human Novel	Novel	Human Novel	Human	Novel	Human Human	Nove1	Human	Nove1 Human	Novel	Nove1	Novel	Nove1	Novel	Novel Human	Novel	Novel Novel Novel
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RRESULT 1
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31-OCT-2002
06-JUN-1991
                                                                                     10-FEB-1986;
14-MAR-1986;
20-OCT-1986;
                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                               Redox active
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                                                                                                                                                                                                                                                                                                                                                                                                     AAP71087;
                                                                                                                                                                                                                                                                                                                                                                                                                           AAP71087 standard; peptide; 6
New and known thioredoxin derivs. and analogues - useful for preventing
                                                                                                                                   10-FEB-1987;
                                                                                                                                                       16-SEP-1987
                                                                                                                                                                            EP237189-A
                                                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                              Unidentified
                      WPI; 1987-258442/37.
                                          Pigiet
                                                                (REPK ) REPLIGEN CORP.
                                                                                                                                                                                                                                                                                                    industry;
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                                                                                                                                                                                                                                                                                                                                   thioredoxin analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Millis CD;
                                                                                                                                                                                                                                                                                                               peptide; antioxidant; stress; ischaemia; lipid peroxidation;
                                                                                                                                                                                                                                                                                                                                                                     (revised)
                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                    86US-00828112.
86US-00839857.
86US-00921287.
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                                                                                                                                                                                                                                                                                                   cosmetics; antibiotic.
                                                                                                                                   87EP-00301150
                                                                                                                                                                                                                      /note= "optional, and may optionally be joined to
terminal blocking gp."
                                                                                                                                                                                                  /note= "see above"
                                                                                                                                                                                                                                                     ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB84815
ADB83930
ADB73085
ADC36923
ADC36923
ADC49944
ADC49143
ADC49521
ADC47521
ADC47521
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ADC1756
ADC1757
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ADD11477
ADD50836
ADD51270
ADD50836
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ADD51350
ADC48897
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ADB78413
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Adc77895
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Adc47266
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Adb87233
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RESULT 2
ABP54936
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           It is a redox active peptide which can be used in cpds. to treat stress and ischaemla, for inhibiting lipid peroxidm. due to drugs or toxic effects, as anti-inflammatory agents, to prevent DNA damage, and to maintain the biological activity of antibiotics or proteins. Thioredoxin can be used as a general antioxidant in the food and cosmetic industry. It may be used in prodm. processes for antibiotics, proteins or polypeptides. Therapeutic dose is 1 microgram/kg - 100mg/kg. (Updated on 11-OCT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct
      glutathionylacetyl)amino)phenylarsenoxide (GSAO) to peptide and protein dithiols. The results indicated that GSAO selectively binds proteins containing closely spaced thiols. The invention provides a method for the treatment and/or prophylaxis of arthritis using a compound of formula A-(L-Y)p, where A is a cell-membrane impermeable pendant group, L is a linker and/or spacer, Y is an arsenoxide or arsenoxide equivalent, and p is an integer from 1 to 10. The compound, e.g. GSAO, is useful for the treatment of arthritic conditions such as calcific periarthritis, enteropathic, chronic, thumb, hip, peripheral, psoriatic, theumatoid, senteropathic, chronic, thumb, hip, peripheral, psoriatic, theumatoid, senteropathic, chronic, thumb, hip, peripheral, psoriatic, theumatoid,
                                                                                                                                                                                                                                              Use of an arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 states.
                                                                                                                                                                      The present sequence is that of a hexapeptide comprising the active site
                                                                                                                                                                                                                                                                                                                                                                                                                    19-MAR-2002; 2002WO-AU000310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          osteopathic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Active site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABP54936;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                metal catalysed oxidative damage in biological reactions and disease
                                                                                                                                                 sequence of thioredoxin. The hexapeptide was us invention that examined the binding of 4-(N-(S-
                                                                                                                                                                                                                 Example 2(b); Page 42; 91pp; English.
                                                                                                                                                                                                                                                                                            WPI; 2002-750519/81.
                                                                                                                                                                                                                                                                                                                                                       (UNIX ) UNISEARCH LTD
                                                                                                                                                                                                                                                                                                                                                                                     19-MAR-2001; 2001AU-00003798
                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200274305-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABP54936 standard; peptide; 6
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or Jaccoud's arthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 WCGPCK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 WCGPCK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page
                                                                                                                                                                                                                                              impermeable cell membrane
in vertebrates.
                                                                                                                                                                                                                                                                                                                         Donoghue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     peptide of thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organoarsenical;
antirheumatic; a:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          arthritis; therapy.
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Pred. No.
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 gout,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                            compound
hand or knee osteoarthritis,
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                                                                                   The present invention relates to a method for producing an antiangiogenesis plasmin fragment. The method comprises contacting a plasminogen polypeptide with a plasminogen activator and a plasmin reductase, where a reduced plasmin protein is produced and the antiangiogenesis plasmin fragment having anti-angiogenesis activity, is released from the reduced plasmin protein. The plasminogen activator is preferably a urokinase-type plasminogen activator. The angiogenesis; collasmin fragment is A61. The plasmin reductase is selected from annexin II heterotetramer, annexin II p36 subunit, p11, thioredoxin, and protein disulphide isomerase. The annexin II heterotetramer is associated with a cell membrane. The method of the invention is useful for modulating (e.g. promoting or inhibiting) angiogenesis. The present sequence represents a mammalian cleavage peptide found in thioredoxin and protein disulphide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anti-angiogenesis plasmin fragment; plasminogen; plasminogen activato: plasmin reductase; reduced plasmin protein; anti-angiogenesis activit; urokinase-type plasminogen activator; A61; annexin II heterotetramer; annexin II p36 subunit; annexin II p11 subunit; thioredoxin; protein disulphide isomerase; modulating angiogenesis; mammalian;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              juvenile osteoarthritis, oligoarthritis, and polyarthritis (all claimed). The compound blocks angiogenesis in the synovial tissue and leucocyte ingress that triggers inflammation, with no signs or symptoms of toxicity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Producing an anti-angiogenesis plasmin fragment, useful for modulating, e.g. promoting or inhibiting angiogenesis, comprises contacting a plasminogen polypeptide with a plasminogen activator and a plasmin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-NOV-2002; 2002US-00304287.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-NOV-2001; 2001US-0333866P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2003-596985/56.
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Pred. No.
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2e+06;
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Sequence 6

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RESULT 4
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PT FOR
                  The specification describes a method for increasing the liquefaction of commucus or sputum. The method comprises contacting the mucus or sputum of the patient with a composition comprises contacting the mucus or sputum of the patient with a composition comprising a protein or peptide containing a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or contacted is located in the respiratory tract, gastrointestinal tract or contacted is located in the respiratory tract, gastrointestinal tract or contacted in a pharmaceutical carrier. The composition is administered to the patient in a pharmaceutical carrier. The protein has a half-life in the companient of about 5 minutes to about 24 hours. The liquid phase of a total coloredox of sample of mucus or sputum from the patient shows a composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin. The composition further comprises nicotinamide-adenine thioredoxin. The composition further comprises nicotinamide-adenine chioredoxin phase trive steep of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Increasing the liquefaction of excessively viscous or cohesive mucus sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mucus; sputum; thioredoxin; active site; cohesive mucus; cohesive sputum; lung disease; cystic fibrosis.
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11-APR-2003;
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2003US-0462082P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                              Query Match
                                                                                                       treating a proliferative condition (especially colon or renal cancer) or a condition mediated by thioredoxin/thioredoxin reductase in the human or animal body. (I) are also useful for inhibiting thioredoxin/thioredoxin reductase and cell cycle progression, to regulate/inhibit cell proliferation and/or to promote apoptosis in a cell in vitro or in vivo. The present sequence represents a thioredoxin family conserved catalytic site peptide, which is given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               treating lung diseases (i.e. cystic fibrosis) in which abnormal or excessive viscosity or cohesiveness of mucus or sputum is a symptom or cause of the disease. The present sequence represents an active site of a thioredoxin which is used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4-(1(Sulfonyl)-1H-indol-2-yl)-4(hydroxy)-cyclohexa-2; 5-dienone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADP87509
                                                                                                                                                                                                                     cyclohexa-2,5-dienone derivatives (I) and their salts, esters, amides, solvates, hydrates or protected forms. (I) have cytostatic activities, and can be used as thioredoxin reductase inhibitors. (I) can be used for
                                                                                                                                                                                                                                                                                                                                            New 4-(1(sulfonyl)-1H-indol-2-yl)-4(hydroxy)-cyclohexa-2,5-dienone derivatives are thioredoxin reductase inhibitors useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                          Stevens MFG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2004056361-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            catalytic site peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADP87509 standard;
                                                                                                                                                                                                                                                                   The present invention described 4-(1(Sulfonyl)-1H-indol-2-yl)-4(hydroxy)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-DEC-2002; 2002WO-GB005842.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-DEC-2002; 2002WO-GB005842
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          colon cancer; renal
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CANC-) CANCER RES TECHNOLOGY LTD.
                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        static; thioredoxin reductase inhibitor; proliferative condition; cancer; renal cancer; thioredoxin; thioredoxin reductase; cycle progression; cell proliferation; apoptosis;
                                                                                                                                                                                                                                                                                                                                 or renal cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ш
l Similarity
6; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 WCGPCK 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
   Conservative
                                                                                                                                                                                                                                                                                                 Page 82; 141pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    family conserved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                          Westwell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               peptide; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                100.0%; Score 47;
100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                          Poole TD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    catalytic site peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                 2e+06;
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                                DB 8;
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                              Length 6;
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RESULT 6
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                                                                                                                                    The present invention describes a composition (I) which modulates the activity of a pl1 protein and effects a change in the level of C plasminogen activation by a cell. Also described: (1) making (M1) a CC clonal cell line, which involves isolating a cell, then characterising the activity of a protein produced by the cell or clonal progeny of the C clonal cell line (II) useful in the identification of composition that C (I) has cytostatic activity, and can be used in antisense therapy. (I) (I) has cytostatic activity, and can be used in antisense therapy. (I) (I) has cytostatic activity, and can be used in antisense therapy. (I) (I) has cytostatic activity, and can be used in antisense therapy. (I) (I) to a cell. The cell can be a human cancer cell, chosen from a HT1080 (I) to a cell. The cell can be a human cancer cell, chosen from a HT1080 (I) to a cell. The cell can be a human cancer cell, chosen from a HT1080 (C cancer in a patient e.g., mouse which involves administering (I) to a cancer cell in the patient. (I) is useful for reducing the development of C cancer cell in the patient. (I) is useful for inhibiting the growth of C cancer inhibiting theour cell invasion in a patient, which involves administering the composition that modulates pl1 activity which involves administering the composition that modulates pl1 activity which involves administering the composition that produces a change in pl1 activity to a cell of the clonal cell line relative to a cell of a clonal cell line that had not received the composition, and C cell is not received the composition, and cell clansing in pl1 activity is a change in the level of plasminogen activation concerns common accommone which is used in the protein activation activation and protein activaty. The composition that produces a change in pl1 activation accommon accommone accomm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Composition useful for inhibiting growth of tumor in patient, modulates activity of pl1 protein and effects change in level of plasminogen activation by a cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense therapy; cancer; tumour; tumour growth inhibition; thioredoxin; protein disulfide isomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-487995/46.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-DEC-2002; 2002US-0433140P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-SEP-2004 (first entry)
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Sequence 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; SEQ ID NO 4; 140pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
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                                                                                                                        disulfide isomerase common
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MEDI-) MEDIOMICS LLC.
(WAIS/) WAISMAN D.
                                                                                  present
                                                                                      invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6
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                                                                                                                        sequence, which is
                                                                                                                            used
                                                                                                                            in
                                                                                                                                the exemplification
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Query Match

100.0%;

Score 47;

DB 8

Length

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RESULT 7
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ID AD09
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AAW45003
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Best Local S
Matches 6
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Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a composition which modulates the activity of a pl1 protein and effects a change in the level of plasminogen activation by a cell. The composition is useful for reducing pl1 protein activity in cancer cells. They are also useful for inhibiting tumour growth or metastasis or treating cancer. The present sequence represents the amino acid sequence of the thioredoxin and protein disulphide isomerase common sequence involved with cleavage, formation and reshuffling of disulphide
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 6 AA;
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Immunomodulatory peptide D7233.
                                            27-APR-1998
                                                                                  AAW45003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      treating cancer or reducing pl1 protein activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New composition a change in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-579971/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-NOV-2001; 2001US-0333866P.
26-NOV-2002; 2002US-00304287
13-DEC-2002; 2002US-0433140P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Thioredoxin and protein disulphide isomerase common sequence.
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                                                                                                                      AAW45003 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Waisman DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-DEC-2003; 2003US-00735577
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                                                                                                                                                                                                                                                             1 WCGPCK 6
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                                                                                                                                                                                                                                                                                                        Similarity 6; Conserv
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                                                                                                                                                                                                                                                                                                     100.0%; ilarity 100.0%; Conservative 0
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                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      which modulates the activity of a pl1 protein and effects level of plasminogen activation by a cell useful for or reducing pl1 protein activity in cancer cells.
                                                                                                                      peptide; 12 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; **
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         activation; tumour growth; metastasis;
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                                                                                                                                                                                                                                                                                                                         Score 47;
Pred. No.
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                                                                                                                                                                                                                                                                                                                         2e+06;
                                                                                                                                                                                                                                                                                                                                                DB 8;
                                                                                                                                                                                                                                                                                                                                            Length 6;
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                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                        0;
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AAY09429
ID AAY0
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present peptide is an immunosuppressant or immunostimulator. An immunosuppressant can be used to treat transplant rejection or autoimmune disease, e.g. rheumatoid arthritis, systemic lupus erythematosis, Sjogren's syndrome, scleroderma, mixed connective tissue disease, dermatomyositis, polymyositis, Reiter's syndrome, Behcet's disease, type I diabetes, Hashimoto's thyroiditis, Graves' disease, multiple sclerosis, myasthenia gravis, encephalomyelitis, phemphigus vulgaris, vegetans or foliaceus, Senear-Usher syndrome or Brazilian phemphigus. An immunostimulator can be used to treat conditions such as cancer or
                       10-OCT-1997;
                                                             06-OCT-1998;
                                                                                                                                         WO9919347-A1
                                                                                                                                                                                                                                                                                                                                                                   AAY09429;
                                                                                                                                                                                                                                                                                                                                                                                                         AAY09429 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acids encoding cysteine- or methionine-containing peptide(s) which have immuno:stimulatory or immunosuppressive activity - can be to treat, e.g. cancers, infection, auto:immune disease or transplant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunomodulator; immunosuppressant; immunostimulator; treatment;
transplant rejection; autoimmune disease; cancer; infection.
                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 12; Page 155; 183pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   rejection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-APR-1996;
23-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-APR-1997;
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                                                                                                                                                                                                                                                                                  Immunoactive
                                                                                                                                                                                                                                                                                                                           14-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9739023-A1
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                                                                                                                                                                                                                       mmune response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A.
                                                                                                                                                                                                                                                                                  peptide SEQ ID NO:54.
                                                                                                                                                                                                                                                                                                                           (first entry
                                                                                                                                                                                                                     immunomodulation; immunosuppression; immunostimulation;
se; immunoreactive; autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96SE-00001422.
96SE-00003469.
                       97US-00949024
                                                               98WO-SE001801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eriksson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97WO-SE000574
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6
                                                                                                                                                                                                                                                                                                                                                                                                         peptide; 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lindvall M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 47; DB 2
Pred. No. 2.3;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                           8
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0;

<u>,,</u> Gaps

0

Length 12; Indels

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RESULT 10
AAU08670
ID AAU08
XX AAU08
AC AAU08
AC AAU08
AC Thio;
XX Thio;
XX Thio;
XX Cell;
KW cchro;
KW cchro;
KW cchro;
XX Synt;
XX Synt;
XX Synt;
XX Synt;
XX O5-M
PR 05-M
PR 07-N
XX OL;
PR 07-N
XX WPI
XX Curt
XX Curt
XX Nov;
PT diff
PT DISC
PT 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes nucleic acid molecules comprising a coding sequence encoding an immunoreactive peptide and further encoding a protein targeting sequence. The nucleic acid is administered to a patient so that its expression product, an immuno-active peptide, modulates an immune response in a patient. The nucleic acid can also be used to treat cancer, either after surgery to remove a portion of the cancer or after ionizing radiation. A cytokine is also administered in conjunction with the nucleic acid. Cells containing the nucleic acid molecule can also be used for treatment. The immunoactive peptide is immunosuppressive and can be used in patients with autoimmune disease. The present sequence represents a specifically claimed immunoactive peptide from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cellular proliferative disorder; differentiative disorder; cancer; carcinoma; sarcoma; tumour; haemopoetic neoplastic disorder; leukaemia; chronic myelogenous leukaemia; cardiovascular disorder; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 32; Page 52; 104pp;
                                                                                           Novel thioredoxin polypeptides and polynucleotides for treating disorder associated with aberrant cellular proliferation or differentiation, e.g., cancer and to identify modulators for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAU08670 standard; protein; 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic genes encoding immunoreactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bergstrand H,
                                                                                        differentiation,
                                                                                                                                                                                                                                                                                                                                                                                   07-MAR-2000; 2000US-0187447P
                                                                                                                                                                                                                                                                                                                                                                                                                                            05-MAR-2001; 2001WO-US007139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Thioredoxin domain markov model sequence #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ASTR ) ASTRA
                                                                                                                                                                                                                                                                                                                            (MILL-) MILLENIUM PHARM INC.
                                                                                                                                                                                                              2001-589945/66
                                                                                                                                                                                                                                                                        RAJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    \vdash
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6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; ilarity 100.0%; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ğ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22105; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eriksson
                                                                                     e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ħ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lindvall M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 47; DB
Pred. No. 2.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peptides containing cysteine or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Saernstrand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>,</u>
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Disclosure;

Fig 3A; 126pp; English

A 14 kDa molecule designated various Enhancing Factor (ECEF); thioredoxin

names (Eosinophil Cytotoxicity-and ADF) and encoding a polypeptide

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RESULT 11
AAR42822
ID AAR43
X DODO DODO DODO DODO DODO X W
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel human thioredoxin polypeptide, 22105, CC fragments of it and allelic variants. 22105 is useful for identifying a ccompound which modulates the activity or expression of the polypeptide cC and nucleic acid, where the binds to the polypeptide, and the binding is considered by direct detection of a test compound/polypeptide binding, by CC using a competition binding assay or an assay for 22105-mediated signal cC transduction, and the activity of the polypeptide is a redox activity or the ability to modulate protein processing, protein folding and protein CC secretion. Anti-22105 antibody is useful for detecting the presence of CC 22105 in a sample. A modulator of 22105 is useful for treating or preventing a disorder such as cellular proliferative or differentiative CC disorder (e.g. cancers, carcinomas, sarcomas and tumours), haemopoetic CC neoplastic disorder (e.g. leukaemia and chronic myelogenous leukaemia) or CC their combination, or an antisense, a ribozyme, ischaemia, thrombosis, CC hypertension), characterised by aberrant activity or expression of a CC preferably a peptide, a phosphopeptide, a small molecule, an antibody or CC their combination, or an antisense, a triple helix molecule, a 22105 nucleic acid or their combination (many examples of each disorder CC are given in the specification). The present sequence represents a CC thioredoxin domain sequence derived from a hidden markov model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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04-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 52 AA;
                                                                                                                                                                                  Silberstein DS,
                                                                                                                                                                                                                                                     06-APR-1992;
06-JUL-1992;
                                                                                                                                                                                                                                                                                                                                      14-OCT-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                       therapy; thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bosinophil cytotoxicity enhancing factor; ECEF; stimulation; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rECEF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR42822;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR42822 standard; protein;
                                               Example 3; Page 67; 96pp;
                                                                             Shortened Eosinophil Cytotoxicity-Enhancing Factor proteins - having Eosinophil-stimulating activity, for use in diagnosis and therapy.
                                                                                                                                                    WPI; 1993-336837/42.
                                                                                                                                                                                                                                                                                                    06-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                        WO9320107-A1
                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                   (BGHM ) BRIGHAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          84.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 WCGPCK 25
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                                                                                                                                  AAQ49405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first entry)
                                                                                                                                                                                                                                                     92US-00862832
92US-00906842
                                                                                                                                                                                                                                                                                                      93WO-US003310
                                                                                                                                                                                                                     & WOMENS HOSPITAL
                                                                                                                                                                                    Balcewicz-Sablinska
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 47;
100.0%; Pred. No.
                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               80
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 52;
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AAR42821
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Best Local S
Matches
                                                                                   A 14 kDa molecule designated various names (Bosinophil Cytotoxicity-Enhancing Factor (BCEF); thioredoxin and ADF) and encoding a polypeptide of 104 amino acids has been found to support the growth of an EBV transformed B lymphocyte cell line in the manner of IL-1; to induce the expression of IL-2 receptors in a large granular lymphocyte cell line from an ATL patient in the manner of IL-1 and to mediate the growth inhibitory properties of interferon gamma. Recombinant ECEF (ECEF) 84 is a truncated version of the 104 amino acid polypeptide comprising the first 84 amino acids of the molecule. Some abortened ECEF's are useful because the support of the strength of the molecule. Some abortened ECEF's are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of 104 amino acids has been found to support the growth of an EBV transformed B lymphocyte cell line in the manner of II-1; to induce the expression of II-2 receptors in a large granular lymphocyte cell line from an ATL patient in the manner of II-1 and to mediate the growth inhibitory properties of interferon gamma. Recombinant ECEF (rECEP) 80 is a truncated version of the 104 amino acid polypeptide comprising the first 80 amino acids of the molecule. Some shortened ECEP's are useful because they retain eosinophil stimulating activity but lack proinflammatory dithiol reductase activity. In particular they can be used for killing tumour cells. (Updated on 25-MAR-2003 to correct PN
because they retain eosinophil stimulating activity but lack proinflammatory dithiol reductase activity. In particular the used for killing tumour cells. (Updated on 25-MAR-2003 to co
                                                                                                                                                                                                                                                                                                                                                                                                                                             Shortened Eosinophil Cytotoxicity-Enhancing Factor proteins - having Eosinophil-stimulating activity, for use in diagnosis and therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eosinophil cytotoxic: therapy; thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAR-2003
04-MAY-1994
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                                                                                                                                                                                                                                                                                                                                                                                           Example 3; Page 66; 96pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Silberstein DS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-APR-1992;
06-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9320107-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR42821;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-OCT-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BGHM ) BRIGHAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1993-336837/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30 WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ49404
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytotoxicity enhancing factor; ECEF; stimulation; diagnosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ₽
Ä
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92US-00862832.
92US-00906842.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              & WOMENS HOSPITAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein;
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Pred. No.
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      they can be correct PN
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RESULT 13
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Best Local S
Matches 6
         can be used in the treatment of various diseases associated with altered mitochondrial function including diabetes mellitus, Huntington's disease, osteoarthritis, Leber's hereditary optic neuropathy (LHON), mitochondrial encephalopathy lactic acidosis and stroke (MELAS), myoclonic epilepsy ragged red fibre syndrome (MERRF) or cancer. Accordingly, these compositions have neuroprotective, nootropic, antidiabetic, anticonvulsant, antiarthritic, osteopathic, ophthalmological and cytostatic activities. This polypeptide sequence is a human heart mitochondrial protein of the invention.
                                                                                                                                                 for therapeutic intervention in treating a disease associated with altered mitochondrial function. Specifically, it refers to a method for identifying proteins of the human heart mitochondrial proteome that are useful for drug screening assays, as well as therapeutic targets. The present invention describes a method for identifying such proteins that
                                                                                                                                                                                                                                                                                                         Identifying a mitochondrial target for drug screening assays and for treating diseases associated with altered mitochondrial function, comprises detecting a modified polypeptide in a sample and correlating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-APR-2002; 2002US-0372843P
17-JUN-2002; 2002US-0389987P
20-SEP-2002; 2002US-0412418P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Huntington's disease; osteoarthritis;
Leber's hereditary optic neuropathy; LHON;
mitochondrial encephalopathy lactic acidosis and stroke; MELAS;
myoclonic epilepsy ragged red fibre syndrome; MERRF; cancer;
neuroprotective; nootropic; antidiabetic; anticonvulsant; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 84
                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 312; 180pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human heat mitochondrial protein as a therapeutic target SeqID312
                                                                                                                                                                                                                                 This invention relates to novel mitochondrial targets that can be
                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-845369/78
                                                                                                                                                                                                                                                                                                                                                                                                                       Ghogh
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BUCK INST AGE RES
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Pred. No. 11;
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                                                                                                                                                                                                                                                                                                                                                                                                                       Gibson BW,
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                                                                                                                                                                                                                                                                                                                                                                                                                       Taylor SW,
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RESULT 14
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            The present invention describes a method (M1) for producing an oil body CC associated with a recombinant multimeric protein complex (MPC). M1 CC comprises producing in a cell comprising oil bodies a first and second CC comprises producing in a cell comprising oil bodies a first and second CC comprises producing in a cell comprising oil bodies a first and second CC complex with man occlusion body CC (OB) through an OB-targeting-protein capable of associating with OB and CC pl. M1 is useful for producing an oil body associated with a recombinant CC mPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition composition composition control treat chronic obstructive pulmonary disease (CDD), cataracts, CC diabetes, envenomation, bronchiopulmonary disease (CDD), ulcers, GERD intestinal (G1) bleeding, intestinal boxed of associating sepsis, gastro and associating the product and association of the surface area of the control of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Loc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic, vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thiocapsa roseopersicina thioredoxin SEQ ID NO:143.
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                                                                                                                                                                                                                                                                                                                                                                                              Claim 81; Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Del Val G,
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Zaplachinski S,
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                                                                                                                                                                                                                                                                                                                                                                                              243; 362pp;
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Moloney M;
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Pred. No.
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18-MAY-1999;
02-JUN-1999;
                                                                                                                                                                                                                          WPI; 200
N-PSDB;
                                                                                               The present invention provides variants of the Aspergillus oryzae pudisulfide isomerase enzyme. These are capable of reducing disulphide bonds in proteins, which may be the cause of allergies in humans. The proteins can be used to reduce the allergenicity of foods, to treat scleroproteins, in detergents, in food additives and in cosmetics
                                                                                                                                                                                 Protein disulfide isomerase variant having increased reducing properties and decreased redox potential than native proteins, used to reduce allergenicity of allergic proteins in feed, food or cosmetic products.
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                                                                                                                                                               Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                Disulphide
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DB; AAC89030.
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37
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                   1 WCGPCK 6
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                                                                                                                                                                                                                                                                                                                                                                                                                      bond; allergy; protein disulfide isomerase variant; detergent;
ive; cosmetic.
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99DK-00000689.
99US-0137068P.
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                                       Score 47; DB
Pred. No. 13;
D; Mismatches
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disulphide
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ABP60779 ID ABPO

ABP60779 standard; protein; 102

8

RESULT 16

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The present invention describes a method (MI) for producing an oil body CC associated with a recombinant multimeric protein complex (MPC). MI CC comprises producing in a cell comprising oil bodies a first and second crecombinant polypeptide (PI, P2), where PI is capable of associating with CC P2 to form the MPC and associating the complex with an occlusion body CC (OB) through an OB-targeting-protein capable of associating with OB and CPI. MI is useful for producing an oil body associated with a recombinant CC MPC. The oil bodies are further formulated for use in the preparation of CC a food product such as milk or wheat based food product, personal care CC product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition CC diabetes, envenomation, bronchiopulmonary disease (COPD), cataracts, CC malignancies, reperfusion injury, wound healing, sepsis, gastro CC disetinal (GI) bleeding, intestinal bowel disease (IBD), ulcers, GERD CR (Gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to invention
                                                                                                                           Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
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                                                                                                                                                                                                                                                    Sequence 102 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 81; Page 238; 362pp; English.
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05-JUL-2001; 2001US-0302885P
04-DEC-2001; 2001US-00006038
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                                                                                                                                                                                                                                                                                                                   invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rooijen G, Deckers F
Val G, Zaplachinski
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                                                                                                                           6,
                                                                                                                                                                Similarity
                                                            WCGPCK 6
WCGPCK 34
                                                                                                                           Conservative
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Moloney M;
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Pred. No.
                                                                                                                               Mismatches
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13;
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WCGPCK 6

Query Match Best Local ( Matches

Local Similarity les 6; Conser

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Mismatches

Indels

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Score 47; DB Pred. No. 13;

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ú 0;

Length 102;

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CC associated with a recombinant multimeric protein complex (MPC). Mi
CC comprises producing in a cell comprising oil bodies a first and second
CC recombinant polypeptide (PI, P2), where PI is capable of associating with
CC P2 to form the MPC and associating the complex with an occlusion body
CC (OB) through an OB-targeting-protein capable of associating with OB and
CP1. Mi is useful for producing an oil body associated with a recombinant
CC P1 oil bodies are further formulated for use in the preparation of
CC a food product such as milk or wheat based food product, personal care
CC product which reduces the oxidative stress on the surface area of the
CC human body or used to lighten the skin, or a pharmaceutical composition
CC used to treat chronic obstructive pulmonary disease (COPD), cataracts,
CC diabetes, envenomation, bronchiopulmonary disease, psoriasis,
CC distro teaches, reperfusion injury, wound healing, sepsis, gastro
CC (gastro osophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to
CC (abscophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to
CC invention. (Updated on 07-AUG-2003 to correct OS field.)
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Sequence
                                                                                                                                                                                                                                                                                                           The present invention describes a method (M1) for producing associated with a recombinant multimeric protein complex (M
                                                                                                                                                                                                                                                                                                                                                              Claim 81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-508806/54.
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05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
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06-SEP-2002
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Val G,
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102
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Zaplachinski
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Moloney M;
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WCGPCK

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ABP607 18
ABP60 XX AB
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05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
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Del Val G, Zaplachinski S,
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Sequence 102
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 81; Page 230; 362pp; English
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Moloney M;
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Query Match
Best Local Similarity
Matches 6; Conserv

Conservative

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Score 47; DB Pred. No. 13; 0; Mismatches

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Length 102; Indels

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Gaps

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Sequence 102

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ABFOO
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                                                                                                                                                        The present invention describes a method (M1) for producing an oil body CC associated with a recombinant multimeric protein complex (MPC). M1 CC associated with a recombinant multimeric protein complex (MPC). M1 CC comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with CC P2 to form the MPC and associating the complex with an occlusion body CC (OB) through an OB-targeting-protein capable of associating with OB and CC P1. M1 is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of CC afood product such as milk or wheat based food product, personal care CC product which reduces the oxidative stress on the surface area of the CC human body or used to lighten the skin, or a pharmaceutical composition CC used to treat chronic obstructive pulmonary disease (CDPD), cataracts, CC diabetes, envenomation, bronchiopulmonary disease (CDPD), cataracts, CC malignancies, reperfusion injury, wound healing, sepsis, gastro intestinal bowel disease, IBD), ulcers, GERD C (Jastro Composition) intestinal based to ARNASCA to ARNASCA to ARNASCA (TARRASCA) and ARRAGAT to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          producing oil body associated with recombinant multimeric protein complex
e.g. redox proteins and immunoglobulins comprises producing recombinant
polypeptides capable of forming the complex in cells comprising oil
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05-JUL-2001; 2001US-0302885P
04-DEC-2001; 2001US-00006038
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Del Val G,
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                                                              (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 ABP60964 represent sequence given in the exemplification of the present invention
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ophthalmological; antidia
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Zaplachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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Moloney M;
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                       The present invention describes a method (M1) for producing an oil body cassociated with a recombinant multimeric protein complex (MPC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease, psoriasis, malignancies, reperfusion injury, wound healing, sepsis, gastro intestinal (GI) bleeding, intestinal bowel disease (IBD), ulcers, GERD (gastro oesophageal reflux disease). ABN89595 to ABN89593 and ABP60677 to ABN89594 represent sequence given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
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05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
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Del Val G, Zaplachinski
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lachinski S,
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Moloney M;
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Pred. No. 13;
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The present invention describes an isolated polynucleotide (I) of CC Alloiococcus otitidis genomic DNA, which encodes an antigenic protein. CC Alloiococcus otitidis is a Gram-positive bacterium. Also described: (1) CC an isolated polypeptide that is encoded by the polynucleotide (I); (2) an expression vector comprising the novel isolated polynucleotide (I); its CC complement, degenerate variant or fragment; (3) a genetically engineered CC host cell, transfected, transformed or infected with the vector of (2); (4) an antibody specific for the polypeptide of (I); (5) an immunogenic composition comprising the polypeptide of (I); (5) an immunogenic composition comprising the polypeptide of (I); (5) an immunogenic composition vector; (6) a pharmaceutical composition comprising the comprising the polypeptide of (I) and a carrier; (7) a protein chip comprising an array configurate of (I) and a carrier; (7) a protein chip comprising an array configurate of (I), their biological equivalent or fragment; (8) immunising against Alloiococcus otitidis by administering to a host the containing the novel polynucleotide, its degenerate variant or fragment; (8) or the antibody of (4); and (II) producing and/or identifying Alloiococcus otitidis in the biological sample; (10) a kit comprising a container container container container container to produce the polymeptide from the culture. (I) can be used in gene therapy. The colymention can be used for treating and daignosing diseases, drug correning assays and monitoring of effects during drug clinical trials. The polymucleotides are useful for expressing and detecting Alloiococcus contents.
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18-NOV-2002; 2002US-0426742P
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Pred. No. 13;
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05-JUL-2001;
04-DEC-2001;
                     associated with a recombinant multimeric protein complex (MPC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (Pl. P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     otitidis. The present sequence represents an Alloiococcus otitidis antigen protein from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oil body; ophthalmological; antidiabetic; cytostatic; antipsorial vasotropic; vulnerary; antibacterial; immunosuppressive; antiulogicod product; milk; wheat; oxidative stress; cataract; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 102 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-SEP-2002
                                                                                                                                                                                                                                                                                                                                                             The present invention describes a method (M1) for producing an oil body
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Val G,
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Zaplachinski S,
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2001US-0302885P.
2001US-00006038.
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Moloney M;
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   This invention describes a novel nucleic acid representing a synthetic CC Bax gene. The Bax gene of the invention is useful for identifying Bax-CC resistant yeast or fungi, identifying, or obtaining and identifying Bax-CC Candida spp. sequences that are differentially expressed in a pathway CC eventually leading to programmed cell death or identifying inhibitors or inhibitor sequences of Bax induced cell death. The products of the CC inhibitor sequences of Bax induced cell death. The products of the CC invention have cyrostatic, fungicide; immunosuppressive, virucide and CC vasotropic activity and can be used in vaccines or for gene therapy. The creating a medicament for treating, pharmaceutical compositions, CC antisense molecules and antibodies are useful as medicaments or in CC preparing a medicament for treating, preventing and/or alleviating CC diseases associated with yeast or fungi or proliferative disorders, such as cancer, or for preventing apoptosis in certain diseases. The compounds or polypeptides, or the genetically modified organism are useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  malignancies, reperfusion injury, wound healing, sepsis, gastro intestinal (GI) bleeding, intestinal bowel disease (IBD), ulcers, GERD (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 ABP60964 represent sequence given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                       New isolated nucleic acid representing a synthetic BAX-gene, useful as medicament for treating, preventing and/or alleviating yeast or fungal infections or proliferative disorders, or for preventing apoptosis in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-DEC-2000; 2000EP-00870318
04-JAN-2001; 2001EP-00870002
09-JAN-2001; 2001EP-00870003
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preparing a medicament for modifying the endogenic
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DB; ABQ76512.
                                                                                                                                                                                                                                                                                       36; Fig 2; 344pp; English.
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Best Local S
Matches 6
        This invention describes a novel nucleic acid representing a synthetic Bax gene. The Bax gene of the invention is useful for identifying Bax-resistant yeast or fungi, identifying, or obtaining and identifying Candida spp. sequences that are differentially expressed in a pathway eventually leading to programmed cell death or identifying inhibitors or inhibitor sequences of Bax-induced cell death. The products of the invention have cycostatic, fungicide; immunosuppressive, virucide and vasotropic activity and can be used in vaccines or for gene therapy. The isolated nucleic acids, polypeptides, pharmaceutical compositions, antisense molecules and antibodies are useful as medicaments or in preparing a medicament for treating, preventing and/or alleviating diseases associated with yeast or fungi or proliferative disorders, such as cancer, or for preventing apoptosis in certain diseases. The compounds or polypeptides, or the genetically modified organism are useful for a medicament for treating the modified organism are useful for the properties of the compounds or polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-DEC-2000; 2000EP-00870318
04-JAN-2001; 2001EP-00870002
09-JAN-2001; 2001EP-00870003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fungal infections. Apoptosis-related diseases include autoimmune disease, ischaemia, diseases related with viral infections or neurodegenerations. This sequence represents a polypeptide associated with the Bax gene described in the disclosure of the invention
                                                                                                                                                                                                                                                                                                                      New isolated nucleic acid representing a synthetic BAX-gene, useful as medicament for treating, preventing and/or alleviating yeast or fungal infections or proliferative disorders, or for preventing apoptosis in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Saccharomyces cerevisiae
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                                                                                                                                                                                                                                                                           Claim
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                                                                                                                                                                                                                                                                        Fig 1; 344pp; English
a medicament for modifying the endogenic flora of humans
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Pred. No.
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13;
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RESULT 25
RESULT
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Bax gene. The Bax gene of the invention is useful for identifying Bax-
resistant yeast or fungi, identifying, or obtaining and identifying
Candida spp. sequences that are differentially expressed in a pathway
eventually leading to programmed cell death or identifying inhibitors or
inhibitor sequences of Bax-induced cell death. The products of the
invention have cytostatic, fungicide; immunosuppressive, virucide and
vasotropic activity and can be used in vaccines or for gene therapy. The
isolated nucleic acids, polypeptides, pharmaceutical compositions,
antisense molecules and antibodies are useful as medicaments or in
preparing a medicament for treating, preventing and/or alleviating
diseases associated with yeast or fungi or proliferative disorders, such
as cancer, or for preventing appotosis in certain diseases. The compounds
or polypeptides, or the genetically modified organism are useful for
preparing a medicament for modifying the endogenic flora of humans and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated nucleic acid representing a synthetic BAX-gene, useful as medicament for treating, preventing and/or alleviating yeast or fungal infections or proliferative disorders, or for preventing apoptosis in certain diseases.
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04-JAN-2001; 2001EP-00870002
09-JAN-2001; 2001EP-00870003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 36; Fig 2; 344pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          describes a novel nucleic acid representing
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Pred. No.
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មានស្ត្រាស់
          片
The invention relates to constructing (MI) a strain of diploid fungal cc cells in which both alleles of a gene are modified, comprising modifying CC one allele by insertion or replacement by a cassette having an CC expressible selectable marker and modifying other allele by cc recombination, of a promoter replacement fragment with a heterologous promoter. (MI) is useful for constructing a strain of diploid fungal CC cells in which both alleles of a gene are modified. The diploid fungal CC cells having both alleles of a gene are modified. The diploid fungal CC contributes to the virulence and/or pathogenicity of a gene that contributes to the virulence and/or pathogenicity of a fungus, a gene that contributes to the virulence and/or pathogenicity of a fungus, and for identifying a therapeutic agent for treatment of a manmalian CC disease. (MI) is useful for identifying a compound which modulates the cativity of a gene product, preferably enzymatic activity, carbon compound catabolism, biosynthetic, transporter, transcriptional,
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Matches
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                                                                                                                                                                                                                                                                                                                                 Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
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20-FEB-2001; 2001US-00792024.
22-AUG-2001; 2001US-0314050P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis; signal transduction; DNA replication; cell division; growth; proliferation; Candida albicans; fungicide; antifungal.
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Pred. No.
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RESULT 27
ABR41607
ID ABR41
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Best Local
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29-MAR-2001; 2001US-0280067P.
29-MAR-2001; 2001US-0280068P.
16-MAY-2001; 2001US-0291280P.
17-MAY-2001; 2001US-0291280P.
17-MAY-2001; 2001US-0291829P.
19-JUN-2001; 2001US-0291949P.
20-JUN-2001; 2001US-0299776P.
20-JUN-2001; 2001US-0300001P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       translational, signal transduction, DNA replication and cell division activity. The method is useful for identifying a compound having the ability to inhibit growth or proliferation of C. ablicans cells and for treating infection by C. albicans. The present sequence is that of an essential Candida abbicans protein used in the method of the invention. Note: The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied to Derwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer; cell proliferative disorder; autoimmune disorder; inflammatory disorder; infection; hormonal disorder; metabolic disorder; neurological disorder; astrointestinal disorder; transport disorder; connective tissue disorder; drug screening; proteome analysis; gene therapy; antisense therapy; genetyping; transgenic animal; knock in; disease model; toxicological testing; transcript imaging; electron transfer-associated protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; dithp;
cancer; cell p
                                                                                                            Daughtery SC, Dam TC, Peralta CH, David MH, Flores V, Marwaha R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human DITHP electron transfer-associated protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 103 AA;
                                                                         WPI; 2003-129518/12.
N-PSDB; ACC46545.
                                                                                                                                                                                                                                                                                                                                           27-MAR-2002; 2002WO-US010056
                                                                                                                                                                                                                                                                                                                                                                                           WO200297031-A2
                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                                                                                       (INCY-)
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                                                                                                  a CH, David
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                                                                                                                                                                                        INCYTE GENOMICS INC.
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                                                                                                                                                               Jones
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                                                                                                                                                Hillman JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnostic and therapeutic polynucleotide; diagnosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein; 103
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                                                                                                                                                             Tran
                                                                                                                                     Yu JY,
Liu TF,
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Pred. No.
                                                                                                               SA,
Lan
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                                                                                                                         Tuason
Nguyen
A, Chen
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                                                                                                               RY,
                                                                                                                Urashka ME;
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13;
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                                                                                                                         Yap PE, Amshey
Kleefeld Y, Ge
Panzer SR, Hai
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Harris B;
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test compound diagnostic and

human diagnostic and therapeutic polypeptide useful for id compound which specifically binds to a polypeptide encoded ostic and therapeutic polynucleotide, and to induce antiboo

identifying ed by human

Novel

Claim 27; SEQ ID NO 1142; 591pp;

English

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The invention relates to novel human diagnostic and therapeutic polymucleotides designated dithp (ACC46080-ACC46749) and to their encoded proteins (DITHP; ABR41136-ABR41812). The invention also relates to polymucleotide sequences at least 90% identical to the dithp cDNA sequences of the invention; recombinant vectors, host cells and transgenic organisms comprising a dithp nucleic acid sequence; the recombinant production of DITHP proteins; antibodies specific for DITHP proteins; microarrays comprising dithp nucleic acid sequences; methods of detecting dithp nucleotide and protein sequences; methods of compounds which specifically bind a DITHP protein, and methods of assessing the toxicity of test compounds using a dithp hybridisation probe. Dithp nucleic acid sequences and DITHP proteins may be used in the diagnosis of a wide variety of conditions including cancer and other cell prolifers; negal or parasitic infections; hormonal disorders; metabolic disorders; negal or parasitic infections; hormonal disorders; metabolic disorders; cransport
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disorders; indiconnective tissue disorders. They may also be used to screen for modulators of protein activity or gene expression. DITHP proteins can additionally be used in analysis of the proteome of a tissue or cell type and to induce antibodies. The dithp nucleic acids are additionally useful in somatic or germline gene therapy of the disorders mentioned above, as a source of antisense sequences, as a source of probes and primers, in genotyping and identification of individuals, in the generation of transgenic animal models of human disease or knock in humanised animals, in toxicological testing, and in transcript imaging. The present sequence represents a DITHP protein which is associated with electron transfer. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
  WPI; 2004-226856/21
N-PSDB; ADK90668.
                                                                                                                                                                                                                                                                                                                                                                                                       Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                      baker's yeast; THX2; transgenic; abiotic stress response; crop plant; oxidoreductase stress-related protein; ORSRP; environmental stress; salinity; drought; temperature; chemical; pathogen; glutaredoxin; GRX; thioredoxin; THX; Quantitative Trait Locus; QTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Baker's yeast thioredoxin 2 (THX2) protein SeqID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 103 AA;
                                                                              Chardonnens A,
                                                                                                                                                                                     07-AUG-2002; 2002EP-00017671
                                                                                                                                                                                                                                        01-JUL-2003; 2003WO-EP006994
                                                                                                                                                                                                                                                                                                                                                  WO2004018687-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-MAY-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disorders; neurological disorders; gastrointestinal disorders; transport
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26
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                                                                                                                                   BASF PLANT SCI GMBH
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                                                                                 Puzio P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 47;
Pred. No.
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13;
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ADM32933
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genes encoding proteins associated with the abiotic stress response. Specifically, it refers to transgenic crop plants expressing the heterologous oxidoreductase stress-related protein (ORSRP) that confers increased tolerance and/ or resistance to environmental stresses such as salinity, drought, temperature, chemicals or pathogens. The present invention describes the ORSRPs as heat-stable gluttaredoxin (GRX) and/ or thioredoxin (THX) proteins, where expression is under the control of an inducible tissue-specific or developmentally-specific promoter and furthermore it provides antisense oligos and RNA interference molecules to inhibit ORSRP expression. In addition, these nucleic acids are useful as Quantitative Trait Locus (QTL) markers that can map genetic loci associated with environmental stress tolerance. This polypeptide sequence
                                                                      Increasing the liquefaction of excessively viscous or cohesive mucus sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site reduced state.
                                                                                                                                                                                                                                                                                                                                                                                                   Saccharomyces cerevisiae. Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                               mucus; sputum; thioredoxin; active site; cohesive mucus;
lung disease; cystic fibrosis; enzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to novel transgenic plant cells transformed by genes encoding proteins associated with the abiotic stress response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New transgenic plant cell transformed by oxidoreductase stress-related protein (DRSRP) coding nucleic acid, useful for producing a transgenic plant with increased environmental stress tolerance.
                                           Disclosure; Page 64; 69pp; English.
                                                                                                                                                           WPI; 2004-270016/25
                                                                                                                                                                                                                                                      10-SEP-2002;
11-APR-2003;
                                                                                                                                                                                                                                                                                                     10-SEP-2003; 2003WO-US028526
                                                                                                                                                                                                                                                                                                                                     25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                     WO2004024868-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Amino acid sequence of a thioredoxin polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-JUN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADM32933 standard; protein; 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           is a baker's yeast THX protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; SEQ ID NO 14; 140pp; English.
                                                                                                                                                                                                                      (NAJE-) NAT JEWISH MEDICAL & RES CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
nes 6; Conserv
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                                                                                                                                                                                                                                                      2002US-0409960P.
2003US-0462082P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cohesive
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The specification describes a method for increasing the liquefaction of mucus or sputum in a patient that has excessively viscous or cohesive

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RESULT 30
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treating lung diseases (i.e. cystic fibrosis) in which abnormal or excessive viscosity or cohesiveness of mucus or sputum is a symptom or cause of the disease. ADM32930ADM32941 represent thioredoxin polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     dinucleotide phosphate (reduced form) (NADPH) for reducing the thioredoxin active site of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a total volume of a sample of mucus or sputum from the patient shows a statistically significant increase after administration of the composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian thioredoxin. The composition further comprises nicotinamide-adenine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the patient with a composition comprising a protein or peptide containing a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the
A 14 kDa molecule designated various names (Eosinophil Cytotoxicity-Enhancing Factor (ECEF); thioredoxin and ADF) has been found to support the growth of an EBV transformed B lymphocyte cell line in the manner of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mucus or sputum.
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06-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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04-MAY-1994
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                                                                                                                     Example 3; Page 65; 96pp; English.
                                                                                                                                                                      Shortened Eosinophil Cytotoxicity-Enhancing Factor proteins Eosinophil-stimulating activity, for use in diagnosis and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-OCT-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eosinophil cytotoxicity enhancing factor; ECEF; stimulation; diagnosis; therapy; thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        rECEF 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         which may used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                      (BGHM ) BRIGHAM
                                                                                                                                                                                                                                                                                                    1993-336837/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WCGPCK 34
                                                                                                                                                                                                                                                                                                                                                               DS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92US-00862832.
92US-00906842.
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                                                                                                                                                                                                                                                                                                                                                                                                                      & WOMENS HOSPITAL.
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                                                                                                                                                                                                                                                                                                                                                            Balcewicz-Sablinska MK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 103;
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                                                                                                                                                                               therapy
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RESULT 31
AAR47855
ID AAR47
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Query Match
Best Local S
Matches 6
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IL-1; to induce the expression of IL-2 receptors in a large granular lymphocyte cell line from an ATL partient in the manner of IL-1 and to mediate the growth inhibitory properties of interferon gamma. Recombinant ECEF (rECEF) 104 is the full length 104 amino acid polypeptide encoded by this molecule and was produced for comparison studies against shortened versions of the polypeptide. Some shortened ECEF's are useful because they retain eosinophil stimulating activity but lack proinflammatory dithiol reductase activity. In particular they can be used for killing tumour cells. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                   The ADF polypeptide suppresses the release of the pancreatic enzymes amylase and lipase into the blood thereby suppressing pancreatitis. The ADF is also capable of eliminating free radicals which can cause organ damage, as well as refolding denatured proteins which have been denatured by free radicals. The ADF may be isolated from cultured human derived cells and prepared by chemical synthesis or recombinant techniques (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                           Use of a polypeptide having human ADF (thioredoxin) activity - for the prophylaxis or therapeutic treatment of pancreatitis or diseases which accompany pancreatic disorders.
                                                        Sequence 104
                                                                                                                                                                                                    Claim 2; Page 9; 15pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADF; thioredoxin; pancreatitis; amylase; lipase; free radicals; protein folding; denaturing; renaturing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human ADF-polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2003
18-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR47855;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR47855 standard; protein; 104 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 104 AA;
                                                                                                                                                                                                                                                                                       N-PSDB; AAQ55111.
                                                                                                                                                                                                                                                                                                      WPI; 1994-027626/04.
                                                                                                                                                                                                                                                                                                                                  Asano T,
                                                                                                                                                                                                                                                                                                                                                                                         19-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                26-JAN-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                             (AJIN ) AJINOMOTO KK
                                                                                                                                                                                                                                                                                                                                                                                                                     18-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30 WCGPCK 35
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 Similarity 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                Kenmochi T,
  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                          92JP-00161454
                                                                                                                                                                                                                                                                                                                                                                                                                     93EP-00109794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                100.0%;
                                                                                                                                                                                                                                                                                                                                  Isono K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
  0
                Score 47;
Pred. No.
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Pred. No.
    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                  Hirakawa T,
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                             BB
                             2;
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                                                                                                                                                                                                                                                                                                                                  Hamuro
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  0
                             Length 104;
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    Indels
  <u>,</u>
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    Gaps
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RESULT 33
AAG80708
ID AAG80
XX AAG80
XX AAG80
XX IS-MA
AC AAG80
XX IS-MA
XX
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AAR72389
ID AAR72
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prepn. of recombinant human ADF by direct expression in E. coli - for treating inflammation and radiation sickness caused by commercially produced free radical(\mathbf{s}).
                                                                                                                                                                                                S. cerevisiae TRX2 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The DNA sequence encoding human ADF was inserted into a plasmid which was then used to transform E.coli. The E.col was then cultured and the recombinantly produced ADF harvested. Human ADF can be used as a treating agent for inflammation and radiation sickness
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Recombinant human ADF
                                                   JP2001288103-A.
                                                                                                    Saccharomyces
                                                                                                                                                                                                                                                 18-MAR-2002
                                                                                                                                                                                                                                                                                                                                              AAG80708 standard; protein; 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2; Page 7-8; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-MAR-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR72389
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (AJIN ) AJINOMOTO KK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Active-site
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DB; AAQ87817.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inflammation; radiation
                                                                                                                                                  yeast;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ||||||
WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                            (first entry)
                                                                                                                                               thioredoxin; lesion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                    cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93JP-00233361
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sickness; fermentation; recombinant.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                  intracellular redox.
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RESULT 34
ABP60927
ID ABP60
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Best Local Similarity
Matches 6; Conser
                                                                                                                                                                                                                                                                                                                             Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antidicer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel thioredoxin (TRX) composition, stable in solution which contains a yeast TRX derived from Saccharomyces cerevisiae X2180-1A. The TRX composition of the invention can be used for treating and/or preventing lesions and diseases related to intracellular redox conditions. This sequence represents the S. cerevisiae TRX2 protein described in the method of the invention
                                                                                                                              19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                         bronchiopulmonary disease; malignancy; gastro intestinal bleeding; intestinal gastro oesophageal reflux disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A thioredoxin composition stable in solution used for treating preventing lesions and diseases related to intracellular redox
                       WPI; 2002-508806/54.
                                                                                                                                                                                                                                       WO200250289-A1
                                                                                                                                                                                                                                                                Staphylococcus aureus.
                                                                                                                                                                                                                                                                                                                                                                                                               Staphylococcus aureus thioredoxin reductase SEQ ID NO:276.
                                                                                                                                                                                                                                                                                                                                                                                                                                           06-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 14-15; 19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-FEB-2000; 2000JP-00025258.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-MAR-2000; 2000JP-00057204.
                                                                                         (SEMB-)
                                                                                                                                                                                  19-DEC-2001; 2001WO-US050240.
                                                                                                                                                                                                              27-JUN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ORIY ) ORIENTAL YEAST CO LTD
                                                Rooijen G, Deckers H,
Val G, Zaplachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002-117628/16.
DB; ABA97065.
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                                                                                         SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; protein; 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                   Heifetz PB,
Moloney M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 47; DB
Pred. No. 14;
0; Mismatches
                                                                                                                                                                                                                                                                                                                     reperfusion injury; wound healing;
                                                              Briggs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                 Dalmia
                                                                 BK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and/or condition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                   sepsis;
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Producing oil body associated with recombinant multimeric protein complex

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RESULT 35
ABP60803
ID ABP60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a method (MI) for producing an oil body associated with a recombinant multimeric protein complex (MPC). MI comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (PI, P2), where PI is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (COB) through an OB-targeting-protein capable of associating with OB and PI. MI is useful for producing an oil body associated with a recombinant CC product such as milk or wheat based for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the CC product which reduces the oxidative stress on the surface area of the CC man body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, CC diabetes, envenomation, bronchiopulmonary disease, psoriasis, CC malignancies, reperfusion injury, wound healing, sepsis, gastro (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to C (preprise the sequence given in the exemplification of the present composition of injury in the exemplification of the present composition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antisportatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antisleer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; chronic obstructive pulmonary disease; chronic obstructive pulmona
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    e.g. redox proteins and polypeptides capable of bodies.
                                                                                                                    (SEMB-)
                                                                                                                                                                                                         19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
                               Van Rooijen
Del Val G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-SEP-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens thioredoxin SEQ ID NO:152
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                                                                                                                    SEMBIOSYS GENETICS INC. SYNGENTA PARTICIPATIONS
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                            G, Deckers H,
Zaplachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                            Heifetz PB,
Moloney M;
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                                                        Briggs SP,
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05-JUL-2001; 04-DEC-2001;

2000US-00742900 2001US-0302885P 2001US-00006038

SEMBIOSYS GENETICS INC. SYNGENTA PARTICIPATIONS

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19-DEC-2000;

19-DEC-2001; 2001WO-US050240

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RESULT 36
ABP60797
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a method (M1) for producing an oil body associated with a recombinant multimeric protein complex (MPC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease, psoriasis, and alignancies, reperfusion injury, wound healing, sepsis, gastro (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to ABP60964 represent sequence given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bos taurus thioredoxin SEQ ID NO:146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABP60797 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-508806/54
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Pred. No. 14;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC associated with a recombinant multimeric protein complex (MPC). M1
CC comprises producing in a cell comprising oil boddies a first and second
CC recombinant polypeptide (P1, P2), where P1 is capable of associating with
CC P2 to form the MPC and associating the complex with an occlusion body
CC P3 to form the MPC and associating the complex with an occlusion body
CC P4 to form the MPC and associating the complex with an occlusion body
CC P5 to form the MPC and associating the complex with an occlusion body
CC P6 to form the MPC and associating with OB and
CC P7 to form the D0 to the complex with a recombinant
CC MPC. The oil bodies are further formulated for use in the preparation of
CC product which reduces the oxidative stress on the surface area of the
CC product which reduces the oxidative stress on the surface area of the
CC product which reduces the oxidative stress on the surface area of the
CC used to treat chronic obstructive pulmonary disease (COPD), cataracts,
CC diabetes, reperfusion injury, wound healing, sepsis, gastro
CC intestinal (G1) bleeding, intestinal bowel disease (IBD), ulcers, GERD
CC (Gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP66677 to
CR CABPEGGA64 represent segmence criven in the averalification of the recombinant of the process.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                  oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
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                               19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
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Del Val G, Zaplachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ovis aries thioredoxin SEQ ID NO:161.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                         WO200250289-A1
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6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                      2001WO-US050240
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Moloney M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>,,</u>
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Briggs SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 104;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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(OB) through an OB-targeting-protein capable of associating with OB and Pl. MI is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or what based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease, psoriasis, malignancies, reperfusion injury, wound healing, sepsis, gastro intestinal (GI) bleeding, intestinal bowel disease (IBD), ulcers, GERD (gastro oesophageal reflux disease). ABN89593 to ABN89593 and ABP60677 to
                                                                                                                                                                                                                                                                                                                                The present invention describes a method (M1) for producing an oil body associated with a recombinant multimeric protein complex (MPC). It comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SEMB-)
Sequence 104
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Del Val G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 81; Page 249; 362pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-508806/54.
                                                                ABP60964 represent sequence given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEMBIOSYS GENETICS INC. SYNGENTA PARTICIPATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G, Deckers H,
Zaplachinski S,
  A
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Moloney M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A
G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Brigge SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dalmia BK;
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밁 Ś Query Match Best Local Similarity Matches 30 WCGPCK 35 1 WCGPCK 6 6, Conservative 100.0%; 0 Score 47; Pred. No. Mismatches DB ۍ ۲۰ 0 Length 104; 0, Gaps

0

ABP60805 standard; protein; 104 ₽

06-SEP-2002 Macaca mulatta (first entry) thioredoxin SEQ ID NO:154

RESULT 38
ABP60805
ID ABP60
XX ABP60
XX ABP60
XX ABP60
XX ABP60
XX ABP60
XX Maca
XX Mult
KW Oill
KW Vasc
KW Chro
KW Chro
KW Goo
KW Oll
KW OBS
KW Goo
KW GOO chronic obstructive pulmonary disease; envenomation; psoriasis; bronchiopulmonary disease; malignancy; reperfusion injury; wound gastro intestinal bleeding; intestinal bowel disease; ulcer; Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antidicer; food product; milk; wheat; oxidative stress; cataract; diabetes; Macaca mulatta. gastro oesophageal disease healing;

19-DEC-2001; 2001WO-US050240

19-DEC-2000; 2000US-00742900

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RESULT 39
ABP60806
ID ABP60806
XX ABP60
XX D6-SE
XX Mus n
XX Multi
KW oil k
KW vasot
KW chror
KW chror
KW gastr
XX G88
XX Mus n
XX Mus n
XX H
XX O200
PN W0200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a method (M1) for producing an oil body cassociated with a recombinant multimeric protein complex (MPC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (C0) through an OB-targeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant CMPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease, psoriasis, malignancies, reperfusion injury, wound healing, sepsis, gastro (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to ABP60964 represent sequence given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                 food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUL-2001; 2001US-0302885P
04-DEC-2001; 2001US-00006038
                                                                                                                                                                                                                          Multimeric protein; redox protein; thioredoxin; thioredoxin reductase, oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e.g. redox proteins and polypeptides capable of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-508806/54
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Del Val G, Zaplachinski S,
                                                           WO200250289-A1
                                                                                              Mus musculus.
                                                                                                                                                                                                                                                                                                       Mus musculus thioredoxin SEQ ID NO:155
                                                                                                                                                                                                                                                                                                                                             06-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                    ABP60806 standard; protein; 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 81; Page 247; 362pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity nes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30
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SYNGENTA P
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                                                                                                                                  oesophageal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WCGPCK 35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                           (first entry)
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Moloney M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Moloney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 47; DB 5; Length 104; Pred. No. 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>.</u>.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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RESULT 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                      Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immonsuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
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05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
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Del Val G, Zaplachinski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (SEMB-)
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                                                                                                                                                                                                                            Oryctolagus cuniculus thioredoxin SEQ ID NO:158.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 104
                                                                                                                                                                                                                                                                                                                                     ABP60809 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    81; Page 247; 362pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                          30
                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 WCGPCK 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA,
                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; S
100.0%; I
tive 0;
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Moloney M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 47; DB Pred. No. 14; D; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 104;
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with

bronchiopulmonary disease; malignancy; reperi gastro intestinal bleeding; intestinal bowel gastro oesophageal reflux disease.

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RESULT 41
ABP60810
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
       Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antilicer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulnonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-DEC-2000; 2000US-00742900
05-JUL-2001; 2001US-0302885P
04-DEC-2001; 2001US-00006038
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-508806/54.
                                                                                                                                                                     Rattus norvegicus thioredoxin SEQ ID NO:159.
                                                                                                                                                                                                       06-SEP-2002
                                                                                                                                                                                                                                        ABP60810
                                                                                                                                                                                                                                                                         ABP60810 standard; protein; 104 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
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                                                                                                                                                                                                                                                                                                                                                                                           1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                   WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 248; 362pp; English.
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Zaplachinski
                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                      (first entry)
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lachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 47;
100.0%; Pred. No.
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Moloney M;
                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          14;
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RESULT 42
ABP60801
ID ABP60
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Matches
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05-JUL-2001; 2001US-0302885P
04-DEC-2001; 2001US-00006038
                                                                                       Gallus gallus
                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        producing oil body associated with recombinant multimeric protein e.g. redox proteins and immunoglobulins comprises producing recomb polypeptides capable of forming the complex in cells comprising oi bodies.
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Del
                                                                                                                                           ABP60801;
                                                                                                                                                                     ABP60801 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200250289-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rattus norvegicus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-DEC-2001; 2001WO-US050240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SEMB-) SEMBIOSYS GENETICS INC. (SYGN ) SYNGENTA PARTICIPATIONS AG
                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rooijen
Val G,
                                                                                                                                                                                                                                      30 WCGPCK 35
                                                                                                                                                                                                                                                               1 WCGPCK 6
                                                                                                                                                                                                                                                                                          Similarity 6; Conser
                                                                                                                                                                                                                                                                                                                                               104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 248-249; 362pp;
                                                                                                                                                                                                                                                                                        100.0%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G, Deckers H,
Zaplachinski S,
                                                                                                                                                                                                                                                                                                                                               A
                                                                                                                  (first
                                                                                       thioredoxin SEQ ID NO:150.
                                                                                                                                                                     protein; 104 AA
                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Heifetz PB,
Moloney M;
                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                        Score 47; DB Pred. No. 14; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                      BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Briggs
                                                                                                                                                                                                                                                                                                                    5
                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                   Length 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SP,
                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dalmia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BK
                                                                                                                                                                                                                                                                                           <u>,,</u>
                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               complex
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vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sep Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic;

wound healing;

0,

bronchiopulmonary disease;

gastro

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RESULT 43
ABG93115
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                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a method (M1) for producing an oil body CC associated with a recombinant multimeric protein complex (MPC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with CC P2 to form the MPC and associating the complex with an occlusion body CC P3 through an OB-teargeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant CC MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, CC diabetes, envenomation, bronchiopulmonary disease, psoriasis, CC malignancies, reperfusion injury, wound healing, sepsis, gastro intestinal (G1) bleeding, intestinal bowel disease (IBD), ulcers, GERD (ABP60964 represent sequence given in the exemplification of the present
                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Van Rooijen G, Deckers .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gastro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200250289-A1
vasotropic; vaccine; gene therapy; proliferative
apoptosis; fungal; yeast; infection; autoimmune d
                                                                                                                                                                                                                                                                                                                                                                                Sequence 104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-508806/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-DEC-2001; 2001WO-US050240
                                Bax; Bax-resistance;
                                                                S. cerevisiae
                                                                                                 21-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 81; Page 245-246; 362pp;
                                                                                                                                                              ABG93115 standard; protein; 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SYGN ) SYNGENTA PARTICIPATIONS
                                                                                                                                                                                                                                                  30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         intestinal bleeding; intestinal bowel disease; ulcer; oesophageal reflux disease.
                                                                                                                                                                                                                                                                                 1 WCGPCK 6
                                                                                                                                                                                                                                                                                                               Similarity 6; Conserv
                                                                                                                                                                                                                                                  WCGPCK 35
                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                 (first entry)
                                                                BAX-associated protein fragment SEQ ID 188.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Deckers H, F
                                                                                                                                                                                                                                                                                                                               100.0%;
               cytostatic; fungicide; immunosuppressive; virucide; gene therapy; proliferative disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Heifetz PB,
Moloney M;
                                                                                                                                                                                                                                                                                                                   <u>..</u>
                                                                                                                                                                                                                                                                                                                               Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ą
                                                                                                                                                                                                                                                                                                                   Mismatches
 autoimmune disease; ischaemia;
                                                                                                                                                                                                                                                                                                                                                DB 5; Length 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Briggs SP,
                                                                                                                                                                                                                                                                                                                 0
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RESULT 44 ADK90667

ADK90667 standard;

protein;

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ZZZXZXZXZ UXSXZXZXZXZ

baker's yeast; THX1; transgenic; abiotic stress response; crop plant; oxidoreductase stress-related protein; ORSRP; environmental stress; salinity; drought; temperature; chemical; pathogen; glutaredoxin; GRX

Baker's yeast thioredoxin 1 (THX1) protein SeqID 12

(first

entry)

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This invention describes a novel nucleic acid representing a synthetic CC Bax gene. The Bax gene of the invention is useful for identifying Bax-CC resistant yeast or fungi, identifying, or obtaining and identifying CC Candida spp. sequences that are differentially expressed in a pathway CC eventually leading to programmed cell death or identifying inhibitors or inhibitor sequences of Bax-induced cell death. The products of the cinvention have cytostatic, fungicide; immunosuppressive, virucide and CC vasotropic activity and can be used in vaccines or for gene therapy. The ci solated nucleic acids, polypeptides, pharmaceutical compositions, CC antisense molecules and antibodies are useful as medicaments or in CC preparing a medicament for treating, preventing and/or alleviating cC diseases associated with yeast or fungi or proliferative disorders, such as cancer, or for preventing apoptosis in certain diseases. The compounds cor polypeptides, or the genetically modified organism are useful for preparing a medicament for modifying the endogenic flora of humans and other mammals. The vaccine is useful for immunising against yeast or fungal infections. Apoptosis-related diseases include autoimmune disease, ischaemia, diseases related with viral infections or neurodegenerations. This sequence represents a polypeptide associated with the Bax gene constitute of the invention
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                                                    Query Match
Best Local S
                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated nucleic acid representing a synthetic BAX-gene, useful as medicament for treating, preventing and/or alleviating yeast or fungal infections or proliferative disorders, or for preventing apoptosis in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-DEC-2000; 2000EP-00870318
04-JAN-2001; 2001EP-00870002
09-JAN-2001; 2001EP-00870003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 certain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Saccharomyces cerevisiae.
                                                                                                              Sequence 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 36; Fig 1; 344pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                neurodegeneration; cell death
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (JANC ) JANSSEN PHARM NV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2002-667002/71.
DB; ABQ76381.
   1 WCGPCK 6
                                      Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diseases.
                                        Conservative
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                                                       100.0%;
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                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Luyten WHML,
                                                           Score 47;
Pred. No.
                                        Mismatches
                                                         14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reekmans
                                                                             ..
                                        0
                                                                             Length 104;
                                        0
                                        Gaps
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RESULT 45
ADO59801
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to novel transgenic plant cells transformed by genes encoding proteins associated with the abiotic stress response. Specifically, it refers to transgenic crop plants expressing the heterologous oxidoreductase stress-related protein (ORSRP) that confers increased tolerance and/ or resistance to environmental stresses such as salinity, drought, temperature, chemicals or pathogens. The present invention describes the ORSRPs as heat-stable glutaredoxin (GRX) and/ or thioredoxin (THX) proteins, where expression is under the control of an inducible tissue-specific or developmentally-specific promoter and furthermore it provides antisense oligos and RNA interference molecules to inhibit ORSRP expression. In addition, these nucleic acids are useful as Quantitative Trait Locus (QTL) markers that can map genetic loci associated with environmental stress tolerance. This polypeptide sequence
                                                                                                                                                                                                               B. subtilis metabolic pathway gene, trxA, protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New transgenic plant cell transformed by oxidoreductase stress-related protein (ORSRP) coding nucleic acid, useful for producing a transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-AUG-2002; 2002EP-00017671
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUL-2003; 2003WO-EP006994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Saccharomyces cerevisiae
11-SEP-2002; 2002DE-01042433.
                                                                                                                                                                               chip; metabolic pathway; acoA; ahpC; ahpF; citB; clpC; clpP; codY; cspA;
                                                                                                                                                                                                                                                 15-JUL-2004
                                                                                                                                                                                                                                                                                                             ADO59801 standard; protein; 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; SEQ ID NO 12; 140pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plant with increased environmental stress tolerance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB; ADK90666.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chardonnens A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     thioredoxin; THX; Quantitative Trait Locus; QTL
                                                                                                Bacillus subtilis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  is a baker's yeast THX protein of the invention.
                                25-MAR-2004
                                                               DE10242433-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BADI ) BASF PLANT SCI GMBH.
                                                                                                                             ; des; dnaK; eno; glnR; groEL; groL; ibpA; ibpB; katA; katE; lctP; opuAB; phoA; phoD; pstS; purC; purN; pyrB; pyrP; sigB; tnrA; trxA; fermentation monitoring; process control; increased efficiency.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2004-226856/21.
                                                                                                                                                                                                                                                                                                                                                                                                  30
                                                                                                                                                                                                                                                                                                                                                                                                                                   μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                  WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                   WCGPCK 6
                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Puzio P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 47; DB
Pred. No. 14;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8; Length 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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RESULT 46
ADS88157
ID ADS88
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a chip that carries probes for at least one regulated genes involved in the corresponding metabolic pathways in other organisms. The specified genes are acoA, ahpC, ahpF, citB, clpC, clpP, codY, cspA, cspB, des, dnaK, eno, glnR, groEL, groL, ibpA, ibpB, katA, katE, lctP, ldh, opuAB, phoA, phoD, pstS, purC, purN, pyrB, gryP, sigB, tnrA, trxA and ydjF. The specification includes a table identifying the products of these genes. The chips are used to determine the products of these genes. The chips are used to determine the physiological condition of organisms being used in a biological process, especially for monitoring fermentations for production of a protein or low molecular weight compound, but also for laboratory-scale cloning and expression analysis. The method is quicker than gel electrophoretic methods currently used, so allows a more rapid intervention for process control and thus increased efficiency. The present sequence represents the amino acid sequence of a B. subtilis metabolic pathway gene protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Feesche J,
Voigt B;
                                                                                                                                                                                                                                                TNP-alpha; chronic inflammatory disease; rheumatoid arthritis; inflammatory bowel disease; infectious disease; septic shock; bacterial infection; neurological disease; stroke-induced inflammation; neurodegenerative disease; cancer; antiinflammatory; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chip carrying probes for specific genes, useful for rapid monitoring organism status, particularly during fermentation.
                                            26-SEP-2002; 2002EP-00021809
10-FEB-2003; 2003EP-00100274.
                                                                                                                                                                  WO2004035783-A2
                                                                                                                                                                                                  Homo
                                                                                                                                                                                                                                    antirheumatic;
                                                                                                                                                                                                                                                                                                                                                     Human protein of a TNF-alpha signalling pathway protein complex SeqID
                                                                                                                                                                                                                                                                                                                                                                                       18-NOV-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADS88157 standard; protein; 104 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 78; 138pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-271050/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-SEP-2002; 2002DE-01042433
                                                                                               24-SEP-2003; 2003WO-EP050655
                                                                                                                                 29-APR-2004
                                                                                                                                                                                                                                                                                                   protein complex; tumour necrosis factor-alpha signalling pathway;
TNP-alpha; chronic inflammatory disease; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HENKEL KGAA.
UNIV GREIFSWALD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WCGPCK 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; ilarity 100.0%; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Maurer K,
                                                                                                                                                                                                                                    cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Breves R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 47; DB Pred. No. 14; 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                멂
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8; Length 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hecker M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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(CELL-) CELLZOME AG

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RESULT 4
AAP92141
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Best Local S
Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to novel protein complexes of the tumour necrosis factor-alpha (TNF-alpha) sending pathway. Specifically, it refers to methods for preparing these complexes comprising at least two component proteins, as well as screening methods to identify modulators of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New protein complex comprising at least one first and second protein the Tumor Necrosis Factor-alpha(TNF-alpha)-signaling pathway, useful diagnosing or treating inflammation, neurological diseases, infectiou diseases or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bouwmeester T, I
Superti-Furga G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              proteins,
                                           12-JUN-1987;
31-MAY-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                            Human adult T cell leukaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAR-2003
25-JUN-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAP92141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAP92141 standard;
(AJIN ) AJINOMOTO KK
                                                                                                                  10-JUN-1988;
                                                                                                                                                                 18-JAN-1989.
                                                                                                                                                                                                               EP299206-A.
                                                                                                                                                                                                                                                                                 Region
                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                    immunodeficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pathway, which include antibodies, agonists and antagonists thereof.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30 WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human adult T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                           87JP-00146348.
88JP-00134218.
                                                                                                                  88EP-00109311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Huhse B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NO 12; 1980pp;
                                                                                                                                                                                                                                                                               Location/Qualifiers
2. .105
                                                                                                                                                                                                                                                             /note= "Claim 1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kruse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein; 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cell leukaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bauch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                            derived factor; hADF; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ruffner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB
14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           derived
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bauer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Þ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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RESULT 48
AAR42819
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
         manner of IL-1. It was also found to induce the expression of IL-2 receptors in a large granular lymphocyte cell line from an ATL patient in the manner of IL-1. The molecule has been designated various names, these
                                 A 14 kDa molecule was isolated by three separate groups and found t support the growth of an EBV transformed B lymphocyte cell line in manner of IL-1. It was also found to induce the expression of IL-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA encoding the polypeptide was sequenced from a gene bank prepd. from mRNA isolated from ATL-2 cells from patient with adult T leukaemia virus. Vectors contg. the DNA can be used to transform host calls for prodn. of hADF polypeptide. The polypeptide causes differentiation and induces growth of lymphocytes and fibroblasts. The N-terminal Met is optional. Similar peptides with deletions, sustns. and allelic derivs. may also be used. See also AAP94809. (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           used
                                                                                                     Shortened Eosinophil Cytotoxicity-Enhancing Eosinophil-stimulating activity, for use in
                                                                                                                                                                                                                          06-APR-1992;
06-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
04-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page; 24pp;
                                                                                                                                                                             Silberstein DS,
                                                                                                                                                                                                                                                              06-APR-1993;
                                                                                                                                                                                                                                                                                                                                                         therapy; thioredoxin
                                                                                                                                                                                                                                                                                                                                                                   Eosinophil
                                                                                                                                                                                                                                                                                                                                                                                           ECEF/thioredoxin/ADF
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR42819;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR42819 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Recombinant human adult T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yodoi J,
                                                                                Disclosure; Fig
                                                                                                                                                                                                                                                                                     14-OCT-1993
                                                                                                                                                                                                                                                                                                           WO9320107-A1
                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                    (BGHM ) BRIGHAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                      1993-336837/42.
Eosinophil Cytotoxicity-Enhancing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6
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                                                                                                                                           AAQ49402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         treating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tagaya Y,
                                                                                                                                                                                                                                                                                                                                                                   cytotoxicity enhancing factor; ECEF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first entry)
                                                                                                                                                                                                                          92US-00862832
92US-00906842
                                                                                                                                                                                                                                                              93WO-US003310
                                                                                6,
                                                                                                                                                                                                     & WOMENS HOSPITAL.
                                                                                                                                                                            Balcewicz-Sablinska
                                                                             96pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein; 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Maeda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
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                                                                                English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cell leukaemia derived factor polypeptide immuno-deficiency disease etc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 47;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ₽
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Factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB
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                                                                                                      Factor proteins diagnosis and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ٠.
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                                                                                                                                                                                                                                                                                                                                                                     stimulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 105;
(ECEF)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hamuro
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                                                                                                                                                                                                                                                                                                                                                                     diagnosis;
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RESULT 50
AAR47856
ID AAR47
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AAR37700
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Matches
                                                                                                                 Matches
                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                    New human
prepn. of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF. The molecule has also been found to mediate the growth inhibitory properties of interferon gamma. Some shortened ECEF's are useful because they retain eosinophil stimulating activity but lack proinflammatory dithiol reductase activity. In particular they can be used for killing tumour cells. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                              Sequence 105 AA;
                                                                                                                                                                                 This sequence represents human ADF. The DNA encoding this sequence may used in the production of a transgenic mouse. The transgenic mouse contains, in the 5' to 3' direction, a human beta-actin promoter, the human ADF coding gene, a termination codon, a human beta-actin poly(A) signal and a human beta-actin terminator. The transgenic mouse is resistant to stress. It is useful in the research of the mechanism of stress and for the development of anti-inflammatory drugs
                                                                                                                                                                                                                                                                                                                                       N-PSDB; AAQ43433.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JP05130819-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR37700 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 105 AA;
            AAR47856 standard;
                                                                                                                                                                                                                                                                            Claim 3; Page 5; 10pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                      12-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                             28-MAY-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                               12-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                resistant; stress;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human;
                                                                                                                                                                                                                                                                                                                                                                         (AJIN ) AJINOMOTO KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                Local Similarity
nes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                  1993-208254/26.
                                                                   3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADF; transgenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6;
                                                                                          WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WCGPCK 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WCGPCK 6
                                                                    WCGPCK 36
                                                                                                                                                                                                                                                                                                  ADF transgenic mouse - is resistant to stress and is useful antiinflammatory drugs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
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                                                                                                                                                                                                                                                                                                                                                                                                                       91JP-00295618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein; 105 AA.
            protein; 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                anti-inflammatory drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mouse; beta-actin; promoter; terminator;
                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 47;
Pred. No.
                                                                                                                            Score 47;
Pred. No.
             ₿
                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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14;
                                                                                                                                        DB
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                                                                                                                                       Length 105;
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                                                                                                                 Indels
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RESULT 51
AAR72390
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                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                 Sequence 105 AA;
                                                                                                                                                                                                                                                                                                                                                    The ADF polypeptide suppresses the release of the pancreatic enzymes amylase and lipase into the blood thereby suppressing pancreatitis. The ADF is also capable of eliminating free radicals which can cause organ damage, as well as refolding denatured proteins which have been denatured by free radicals. The ADF may be isolated from cultured human derived cells and prepared by chemical synthesis or recombinant techniques (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Use of a polypeptide having human ADF (thioredoxin) activity - for the prophylaxis or therapeutic treatment of pancreatitis or diseases which accompany pancreatic disorders.
                              JP07079780-A
                                                                                                                            Recombinant human
                                                                                                                                                10-NOV-1995
                                                                                                                                                                     AAR72390;
                                                                                                                                                                                        AAR72390 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; Page 12; 15pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Asano T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EP579958-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADF; thioredoxin; pancreatitis; amylase; lipase; free radicals; protein folding; denaturing; renaturing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human ADF-polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
18-JUL-1994
                                                                                  Homo
                                                                                                       ADF; inflammation; radiation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-JAN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR47856;
                                                  ctive-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (AJIN ) AJINOMOTO KK.
                                                                                 sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1994-027626/04.
                                                                                                                                                                                                                                             31
                                                                                                                                                                                                                                                                  \vdash
                                                                                                                                                                                                                                              WCGPCK 36
                                                                                                                                                                                                                                                                  WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kenmochi T,
                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   92JP-00161454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93EP-00109794.
                                                  Location/Qualifiers 32. .35
                                                                                                                            ADF.
                                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isono
                                                                                                                                                                                           105 AA.
                                                                                                                                                                                                                                                                                       0,
                                                                                                       sickness;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ~
                                                                                                                                                                                                                                                                                                   Score 47; DB Pred. No. 14;
                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hirakawa
                                                                                                       fermentation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hamuro
                                                                                                                                                                                                                                                                                                            Length 105;
                                                                                                                                                                                                                                                                                       Indels
                                                                                                        recombinant
                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                       0;
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28-MAR-1995.

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RESULT 52
AAW69567
ID AAW69
XX AAW69
AC AAW69
XX Human
XX
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Best Local S
Matches 6
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produced
New thioredoxin variants with amino acid substitutions for cysteine except at active centre for increasing stability in non-reducing conditions - used with Ref-1 to enhance AP-1 transcriptional activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The DNA sequence encoding human then used to transform E.coli. I recombinantly produced ADF harve agent for inflammation and radia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW69567 standard; protein; 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB;
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                                                                                         WPI; 1998-364621/32.
                                                                                                                                                                                                                                             14-JAN-1997;
                                                                                                                                                                                                                                                                                           14-JAN-1998;
                                                                                                                                                                                                                                                                                                                                       15-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                              EP853088-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW69567;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (AJIN
                                                                                                                                                                                                   (ORIY)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity les 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1995-157850/21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        thioredoxin mutant C62S/C69S/C73S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of recombinant human ADF by direct expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            thioredoxin; mutant; variant; TRX; AP-1; transcriptional activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AJINOMOTO
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                                                                                                                                                                            ORIENTAL YEAST CO LTD YODOI J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inflammation and radiation sickness caused by commercially free radical (s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WCGPCK 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                               97JP-00004489.
                                                                                                                                                                                                                                                                                           98EP-00100535
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers 62
                                                                                                                                                                                                                                                                                                                                                                                                                            /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ing human ADF was inserted into a plasmid which was E.coli. The E.coli was then cultured and the ADF harvested. Human ADF can be used as a treating and radiation sickness
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Сув
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Сув
                                                                                                                                                                                                                                                                                                                                                                                                                            gy.
                                                                                                                                                                                                                                                                                                                                                                                                                            in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in wild type"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in wild
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                            wild
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                                                                                                                                                                                                                                                                                                                                                                                                                            type"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             type"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB
14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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                                                                        Query Match
Best Local
                                                            Matches
                                                                                                                                                                    The present sequence represents a human thioredoxin (TRX) variant derived from the wild-type sequence (obtained from Genbank), as stated in the specification. TRX variants can be made stable under non-reducing conditions by substituting at least one or all Cys residues by other amino acid residues, except that Cys residues in the active centre of TRX remain unmodified. Also described in the present invention is a factor for enhancing the transcriptional activity of AP-1 which comprises the TRX variants can be used with Ref-1 to enhance AP-1 transcriptional activity. TRX variants do not form multimers and are stable under non-reducing
                                                                                                                           Sequence 105
                                                                                                                                                                                                                                                                                                                                              Example
                                                                                                                                                          conditions
31 WCGPCK 36
                               1 WCGPCK 6
                                                           6; Conserv
                                                                                                                                                                                                                                                                                                                                             3; Page; 14pp;
                                                            Conservative
                                                                                                                           AA;
                                                                           100
                                                                                                                                                                                                                                                                                                                                              English
                                                                             .0%;
                                                              0,
                                                           Score 47; DB
Pred. No. 14;
); Mismatches
                                                                                           DB
                                                                                           2
•-
                                                              0
                                                                                           Length 105;
                                                              Indels
                                                              0
                                                              Gaps
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AAY25908
                                                                           RESULT 53
                    06-OCT-1999
                                      AAY25908;
                                                         AAY25908
Human thioredoxin protein.
                                                         standard;
                   (first
                                                         protein;
                   entry)
                                                          105
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Thioredoxin; thioredoxin reductase; l cytostatic; tumour growth inhibitor; phosphorothioate linkage. Homo sapiens reductase; human; antisense; primer; metastasis; h inhibitor; detection; nuclease resistant;

WO9938963-A1

05-AUG-1999.

29-JAN-1999;

99WO-CA000077

30-JAN-1998; 98US-0073196P

(GENE-) GENESENSE TECHNOLOGIES INC.

JA, Young Ħ, SA

WPI; 1999-469328/39. AAZ00544.

Antisense oligonucleotides against thioredoxin and thioredoxin reductase genes, useful for inhibiting tumor growth and metastasis.

Disclosure; Fig 2; 88pp; English.

This invention describes novel antisense oligonucleotides against thioredoxin and thioredoxin reductase gene which have cytostatic activity and are useful for inhibiting tumour growth and metastasis in mammals. They may also be used as hybridization probes to detect the presence of the thioredoxin and thioredoxin reductase mRNAs in mammalian cells. They may also be used as molecular weight markers. The antisense oligonucleotides are nuclease resistant due to the presence of phosphorothioate internucleotide protein linkages. This sequence represents

Sequence 105 A

<u>ب</u>

Length 105

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RESULT 54
AAG03956
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                                                Query Match
Best Local S
Matches 6
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Best Local Similarity
Matches 6; Conserv
                                                                                                                  prepared from mRNAs encoding secreted proteins. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 3' untranslated region (UTR) of the mRNA because they are often obtained from 30 different primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion vectors
                                                                                                  Sequence 105 AA;
                                                                                                                                                                                                                                                                      The present sequence is a polypeptide encoded by one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. The 5' ESTs warm
                                                                                                                                                                                                                                                                                                          Claim 13; SEQ ID NO 8037; 71pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                    diagnostic, forensic,
                                                                                                                                                                                                                                                                                                                                               New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo
                                                                                                                                                                                                                                                                                                                                                                                                                             Dumas Milne Edwards
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-FEB-2000; 2000EP-00200610
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GEST ) GENSET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy;
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                                               Similarity 6; Conserv
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WCGPCK 36
                    WCGPCK 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EST;
                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chromosome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein,
                                                           100.0%;
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                                                                                                                                                                                                                                                                                                                                   gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                             Duclert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence tag; secreted protein; cDNA isolation; mapping.
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                                               Score 47; DB
Pred. No. 14;
D; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 47;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                            Giordano
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. 14;
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                                                                          DB
                                                                       3; Length 105;
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                                                Indels
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                                           Gaps
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RESULT 55 ABP60700

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31

WCGPCK 36

1 WCGPCK 6

0

Indels

0

Gaps

0;

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The present invention describes a method (M1) for producing an oil body CC associated with a recombinant multimeric protein complex (MPC). M1 CC comprises producing in a cell comprising oil bodies a first and second CC recombinant polypeptide (P1, P2), where P1 is capable of associating with CC P2 to form the MPC and associating the complex with an occlusion body CC (OB) through an OB-targeting-protein capable of associating with OB and CC P1. M1 is useful for producing an oil body associated with a recombinant CC MPC. The oil bodies are further formulated for use in the preparation of CC a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the CC muman body or used to lighten the skin, or a pharmaceutical composition CC used to treat chronic obstructive pulmonary disease (COPD), cataracts, CC diabetes, envenomation, bronchiopulmonary disease (COPD), cataracts, CC malignancies, reperfusion injury, wound healing, sepsis, gastro CC (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to ABP60964 represent sequence given in the exemplification of the present
Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-DEC-2000;
05-JUL-2001;
04-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil bodies.
                                                                                                                                        Sequence 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 203; 362pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N-PSDB; ABN89591.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-JUN-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human thioredoxin protein SEQ ID NO:47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-SEP-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-508806/54.
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Val G,
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100.0%; Score 47; DB ilarity 100.0%; Pred. No. 14; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G, Deckers H,
Zaplachinski S,
                                                                                                                                               B
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2001US-0302885P.
2001US-00006038.
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Moloney M;
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                                                                         Length 105;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            묤,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sepsis;
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RESULT 56
ABP60757
                                                                       The present invention describes a method (M1) for producing an oil body case of the with a recombinant multimeric protein complex (MPC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease (COPD), cataracts, analignancies, reperfusion injury, wound healing, sepsis, gastro intestinal (GI) bleeding, intestinal bowel disease (IBD), ulcers, GERD (gastro oesophageal reflux disease). ABN89559 to ABN89593 and ABP60677 to ABP60964 represent sequence given in the exemplification of the present
              Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Van Rooijen
Del Val G,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-SEP-2002
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                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 81; Page 230; 362pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcy
food product; milk; wheat; oxidative stress; cataract; diabetes;
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                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SYNGENTA PARTICIPATIONS
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                                                                       105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      c protein; redox protein; thioredoxin; thioredoxin reductase;
ophthalmological; antidiabetic; cytostatic; antipsoriatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zaplachinski
   Conservative
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                100.0%;
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Moloney M;
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                Score 47;
Pred. No.
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   Mismatches
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                                  Length 105
   Indels
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Query Match
Best Local Similarity
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                                                                                                                      preparing a medicament for treating, preventing and/or alleviating diseases associated with yeast or fungi or proliferative disorders, such as cancer, or for preventing apportists in certain diseases. The compounds or polypeptides, or the genetically modified organism are useful for preparing a medicament for modifying the endogenic flora of humans and other mammals. The vaccine is useful for immunising against yeast or fungal infections. Apoptosis-related diseases include autoimmune disease, ischaemia, diseases related with viral infections or neurodegenerations. This sequence represents a polypeptide associated with the Bax gene described in the disclosure of the invention
                                                                                                                                                                                                                                                                                                                                                         This invention describes a novel nucleic acid representing a synthetic Bax gene. The Bax gene of the invention is useful for identifying Bax-resistant yeast or fungi, identifying, or obtaining and identifying Candida spp. sequences that are differentially expressed in a pathway eventually leading to programmed cell death or identifying inhibitors inhibitor sequences of Bax-induced cell death. The products of the invention have cytostatic, fungicide; immunosuppressive, viricide and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid representing a synthetic BAX-gene, useful as medicament for treating, preventing and/or alleviating yeast or fungal infections or proliferative disorders, or for preventing apoptosis in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-DEC-2000; 2000EP-00870318
04-JAN-2001; 2001EP-00870002
09-JAN-2001; 2001EP-00870003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             vasotropic; vaccine; gene therapy; proliferative disorder; cancer; apoptosis; fungal; yeast; infection; autoimmune disease; ischaemia; neurodegeneration; cell death.
                                                                                      Sequence 105
                                                                                                                                                                                                                                                                                                       isolated nucleic acids, polypeptides, pharmaceutical compositions, antisense molecules and antibodies are useful as medicaments or in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human BAX-associated protein fragment SEQ ID
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                                                                                                                                                                                                                                                                                                                                        vasotropic activity and can be used in vaccines or for gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002-667002/71.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          36; Fig 2; 344pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ungicide; immunosuppressive; virucide; proliferative disorder; cancer;
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Conservative

100.0%;

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Mismatches

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WCGPCK 36 WCGPCK 6

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The invention relates to identifying (M1) a candidate p53 pathway combiling agent, by contacting an assay system comprising a purified HM polypeptide (human orthologue of genes that modify the p53 pathway in CC prosophila) or nucleic acid with a test agent under conditions, where but for the presence of the test agent, the system provides a reference cativity, and detecting a test agent, the system provides a reference cativity, and detecting a test agent biased activity of the assay system. CC Also included are modulating (M2) a p53 pathway of a cell (comprising contacting a cell defective in p53 function with a candidate modulator that specifically binds to a HM polypeptide comprising (M3) a p53 pathway of a cell (comprising contacting the cell with an agent that specifically binds an HM polypeptide or mucleic acid) and diagnosing (M4) a disease in a patient (comprising contacting the cell with an agent that specifically binds an HM polypeptide or mucleic acid) and diagnosing (M4) a disease in a patient (comprising: (a) obtaining a biological sample compression; (c) comparing the results with a control; and (d) determining (M3) a p53 pathway (d) contacting the sample with a probe for HM compression; (d) comparing the results with a control; and (d) determining (M3) assetul for diagnosing breast, colon, kidney, lung and ovarian cancer, in a patient, where the cancer has greater than 25 % expression level.

CC the p53 pathway, such as, angiogenesis, apoptotic or cell to defects in the p53 pathway, and as angiogenesis, apoptotic or cell composition or the cell, so that the cell undergoes normal control for treating defects in the p53 pathway such as angiogenic, also useful for treating defects in the p53 pathway such as angiogenic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 58
ABO07225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying modulators of the p53 pathway for use in treating apoptotic or cell proliferation disorders, comprises screening for agents that modulate activity of a human ortholog of genes that modify the p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUN-2001; 2001US-0296076F
10-OCT-2001; 2001US-0328605F
15-FEB-2002; 2002US-0357253F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; p53 modifier; cytostatic; cancer; cytostatic; antiangiogenic; antiapoptotic; p53 pathway; breast cancer; colon cancer; kidney cancer; lung cancer; ovarian cancer; andiogenesis; cell cycle; apoptotic disorder; cell proliferation disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 536; 678pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pathway in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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RESULT 59
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Best Local Similarity
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                                     ACF12828 to ACF12947 encode the human cervical cancer marker proteins (I) given in ARB92047 to ABR92164. A higher level of expression of (I) than normal indicates the presence of cervical cancer. Also described: (1) a vector (II) containing (I); (2) a host cell (III) containing (I); and (3) assessing (MI) whether a patient is afflicted with cervical cancer, comprising comparing the level of expression of a marker in a patient's sample, and the normal level of expression of the marker in a control non-cervical cancer sample, where a significant increase in the level of expression of the marker in the patient's sample relative to that in the control sample is an indication that the patient is afflicted with cervical cancer. (I) has cytostatic activity, and can be used in gene therapy and in vaccines (I) is useful in detecting, characterising, preventing and treating human cervical cancers. (I) may also be used in various prognostic and diagnostic assays, pharmacogenomics and in
                                                                                                                                                                                                                                                                                                              New isolated nucleic acid molecule useful for detecting, characterizing, preventing and treating human cervical cancers, in various prognostic an diagnostic assays, in pharmacogenomics and in monitoring clinical trials
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 105 AA;
                                                                                                                                                                                                                                                                                   Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; ACF12941.
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13-JUN-2001; 2001US-0298159P
14-NOV-2001; 2001US-0335936P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human cervical cancer cell marker protein SEQ ID NO:226
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  detection; gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6;
                                                                                                                                                                                                                                                                                  Page 377-378; 386pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WCGPCK 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                Chen Y,
M, Glatt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cell proliferation disorders. The present sequence human p53 pathway modifying protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer; cervical cancer marker; cancer therapy;
therapy; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Monahan JE,
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                                                                                                                                                                                                                                                                                                                 in monitoring clinical trials.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Kamatkar
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Query Match Best Local Similarity

100.0%;

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No. DB 14;

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ADD67269 ID ADD6

RESULT

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Matches

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comprises: (a) capture agents, where each capture agent is preselected to specifically bind to a pre-selected tags, and (b) tagged reagents, each comprising one of the pre-selected tags, where each locus in the collection comprises the same capture agent, where the tagged reagent comprises a molecule and a tag, each tag is bound to a capture agent, where cach tag is bound to a capture agent, where each tag is bound to a capture agent, where each locus comprises tagged reagent with the capture agent, where each locus comprises tagged reagents and where each of the different molecules at each locus comprises the same pre-selected tag. Also described: (1) a system for profiling samples; (2) a method for profiling a sample; (3) a computer system or computer readable medium comprising the database produced by the method of profiling a sample; (4) a method for proparing a capture system that displays a collection of binding sites comprising capture agents bound to a solid support and tagged reagents; and (6) a method for screening samples. The combination (1) is useful for some property agents in the scenario for the capture agents and tagged reagents.
                                                                                                                 Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New combination comprising an addressable collection of binding sites, software comprising instructions for pattern recognition and an imager for detecting patterns, useful for profiling a sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 apparatus combination; binding site collection; pattern recognition; profiling; screening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        comprising: (a) an addressable collection of binding sites; and (b) software comprising instructions for pattern recognition and/or an imager for detecting patterns. The addressable collection of binding sites comprises: (a) capture agents, where each capture agent is preselected to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADD67269;
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                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                           profiling a sample. The present of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a combination of apparatus (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 7; SEQ ID NO 127; 309pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-636736/60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (POIN-) POINTILLISTE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-JAN-2002; 2002US-0352011P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-JAN-2003; 2003WO-US002397
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JAN-2004
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                                                                                   Similarity 6; Conserv
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WCGPCK 6
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                                                                                                                    100.0%;
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                                                                                         <u>,</u>
                                                                                                                    Score 47; DB
Pred. No. 14;
                                                                                                                                                                                                                                                                                         sequence is
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                                                                                      Mismatches
                                                                                                                                               DB 7;
                                                                                                                                                                                                                                                                                         used in the exemplification
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ADF30728:
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XX COTO
XX RAT
XX US20
PN US20
XX US
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ADJ70213
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DT 06-M2
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KW Mitoc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a method of treating an angiogenesis-mediated disorder in a subject. The method is useful for treating angiogenesis-mediated disorder, e.g., retinal or choroidal neovascularisation or diseases associated with chronic inflammation, myocardial ischaemia, stroke, coronary artery disease or peripheral vascular disease. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Treating angiogenesis-mediated disorder, e.g., retinal or choroidal neovascularization or diseases associated with chronic inflammation, myocardial ischemia, stroke, coronary artery disease or peripheral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rat; angiogenesis; angiogenesis modulating protein; retinal neovascularisation; choroidal neovascularisation; chronic inflammation; myocardial ischaemia; stroke;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-FEB-2002; 2002US-0355295P
26-JUN-2002; 2002US-0391758P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; SEQ ID NO 291; 26pp; English
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N-PSDB; ADF30727.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vascular disease.
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  Huntington's disease; osteoarthritis;
                                mitochondrial; human;
                                                                                         Human heat mitochondrial protein as a
                                                                                                                                                  06-MAY-2004
                                                                                                                                                                                                                                                                 ADJ70213 standard; protein; 105 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PROC ) PROCTER & GAMBLE CO.
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les 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002US-00316253
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100.0%; Pr
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                                screening assay; diabetes mellitus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein #146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 47;
Pred. No.
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                                                                                         therapeutic target SeqID2019.
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                                                                                                                                                      RESULT 63
                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                 for therapeutic intervention in treating a disease associated with altered mitochondrial function. Specifically, it refers to a method for identifying proteins of the human heart mitochondrial proteome that are useful for drug screening assays, as well as therapeutic targets. The present invention describes a method for identifying such proteins that can be used in the treatment of various diseases associated with altered mitochondrial function including diabetes mellitus, Huntington's disease, osteoarthritis, Leber's hereditary optic neuropathy (LHON), mitochondrial encephalopathy lactic acidosis and stroke (MELAS), myoclonic epilepsy ragged red fibre syndrome (MERRF) or cancer. Accordingly, these compositions have neuroprotective, nootropic, antidiabetic, antignant antiarthritic corporation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Leber's hereditary optic neuropathy; LHON; mitochondrial encephalopathy lactic acidosis and stroke; MELAS; myoclonic epilepsy ragged red fibre syndrome; MERRF; cancer; neuroprotective; nootropic; antidiabetic; anticonvulsant; antiarthritic; osteopathic; ophthalmological; cytostatic.
Antidiabetic; Antiarteriosclerotic; Neuroprotective; Nootropic; Antiparkinsonian; Anticonvulsant; Cardiant; Cardiovascular; Antirheumatic; Antiarthritic; Cytostatic; Antipsoriatic; Gene T
                                                                                                                                                                                                                                                                                           Sequence 105
                                                                                                                                                                                                                                                                                                                 anticonvulsant, antiarthritic, osteopathic, ophthalmological and cytostatic activities. This polypeptide sequence is a human hear mitochondrial protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying a mitochondrial target for drug screening assays and treating diseases associated with altered mitochondrial function, comprises detecting a modified polypeptide in a sample and correl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ghosh SS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-APR-2002; 2002US-0372843P.
17-JUN-2002; 2002US-0389987P.
20-SEP-2002; 2002US-0412418P.
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                                                                            06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention relates to novel mitochondrial targets that can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003087768-A2
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                                                                                                                           ADJ47593 standard; protein; 105
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(BUCK-) BUCK INST AGE RES.
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                                                  coding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2019; 180pp; English.
                                                                            entry)
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                                                  sequence,
                                                                                                                                                                                                                                          100.0%; Score 47; Di
100.0%; Pred. No. 14
tive 0; Mismatches
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                                                  SEQ ID 1.
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                                                                                                                                                                                                                                                     DB 7;
14;
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Gene Therapy;
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BE Human
XX ITHMUM
XW ANTI-S
KW Gene
KW ITHMUM
XX Gene
KW ITHMUM
XX
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                                                                                                                                                                                                                                                                              Query Match
Best Local
                       Immunosuppressive;
Antiallergic; Muscu
Gene Therapy; PRO;
                                                                                                     17-JUN-2004
                                                                                                                                                                                                                                                                                                                    Sequence 105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Min W,
                                                                            Human
                                                                                                                             ADL82895;
                                                                                                                                                    ADL82895 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2004013283-A2
               lmmune-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYRP ) UNIV ROCHESTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2004-157111/15.
DB; ADJ47596.
                                                                                                                                                                                                                    31
                                                                                                                                                                                                                                           1 WCGPCK 6
                                                                                                                                                                                                                                                                   6;
                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                  WCGPCK 36
                                                                                                                                                                                                                                                                   Conservative
                                                                                                     (first
                         sive; Cytostatic; Antiarthritic; Antirheumatic; Antianemic;
Muscular; Neuroprotective; Nephrotropic; Antiinflammatory;
PRO; B cell related disorder; cancer;
                                                                            SEQ
                                                                                                                                                   protein;
                                                                            IJ
                                                                                                    entry)
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diabetes, apoptotic disease, mitochondrial dysfunction disease or cardiac dysfunction. Mitochondrial dysfunction diseases include Kearns-Sayre syndrome or encephalomyopathy lactic acidosis and stroke-like episodes. Apoptotic diseases include neurodegenerative diseases, e.g. Allheimer's disease, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, multiple sclerosis or peripheral neuropathy. Cardiac dysfunction includes myocardial infarction, cardiomyopathy, arterial hypertension or heart failure. The mutant Trx proteins are also useful for treating or diagnosing angiogenesis dependent diseases, e.g. rheumatoid arthritis, cancer or psoriassis. The present sequence is the wild-type Trx protein, from which the mutants of the invention were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the oxidizing effects of cytokines or reactive oxygen species or resistant to S-nitrosylation of a SH-group by nitrous oxide. The multiple proteins are useful for treating or diagnosing atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New mutant thioredoxin molecule which is resistant to oxidizing effects of cytokines or reactive oxygen species, useful for treating or diagnosing atherosclerosis, diabetes, Alzheimer's disease, myocardial infarction or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 thioredoxin; Trx; atherosclerosis; diabetes; apoptotic disease; mitochondrial disease; cardiac dysfunction; angiogenesis dependent disease; neurodegenerative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to mutant thioredoxin (Trx) proteins and their coding sequences, where the thioredoxin molecule is resistant to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 1; 145pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-AUG-2002; 2002US-0401073P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-JUL-2003; 2003WO-US022847.
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                                  100.0%;
0,
Score 47; DB
Pred. No. 14;
0; Mismatches
                                  DB
14;
                                                                             8
0
                                                                        Length 105
<u>.</u>
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Gaps
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RESULT 65
ADM32938
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              deficiency, selective deficiency of IgG subclasses, immunodeficiency with hyper IgM, transient hypogammaglobulinemia of infancy, Burkitt's lymphoma, intermediate lymphoma, follicular lymphoma, type II hypersensitivity, rheumatoid arthritis, autoimmune mediated haemolytic anaemia, myaethenia gravis, hypoadrenocorticism, Jlomerulonephritis, or ankylosing spondylitis. The PRO proteins are also useful for preparing a medicament for treating a condition that is responsive to the PRO protein, e.g. cancer or immune-mediated inflammatory diseases. The PRO coding sequences are useful as hybridization probes in chromosome and gene mapping, in preparing PRO proteins, or in generating transgenic animals or knockout animals, which in turn are useful in the development and screening of therapeutically useful reagents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New PRO polypeptide, useful for diagnosing and treating a B cell related disorder, e.g. Burkitt's lymphoma, rheumatoid arthritis, autoimmune mediated hemolytic anemia, myasthenia gravis or ankylosing spondylitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 10; Fig 97; 695pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; ADL82894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-329389/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chiu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-SEP-2002; 2002US-0411392P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                            mucus; sputum; thioredoxin; active site; cohesive mucus; cohesive sputum; lung disease; cystic fibrosis; enzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to PRO proteins and their coding sequences. The PRO proteins are useful for diagnosing and treating a B cell related disorder, e.g. X-linked infantile hypogammaglobulinemia, polyeaccharide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-SEP-2003; 2003WO-US029097
10-SEP-2003; 2003WO-US028526.
                                 25-MAR-2004
                                                                                                                                                                                                                               17-JUN-2004
                                                                                                                                                                                                                                                                                            ADM32938 standard; protein; 105 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antigen unresponsiveness, selective IgA deficiency, selective IgM deficiency, selective deficiency of IgG subclasses, immunodeficiency with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wu TD;
                                                                WO2004024868-A2
                                                                                                             Homo sapiens
                                                                                                                                                                                               Amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GETH )
                                                                                                                                                                                                                                                                                                                                                                             31
                                                                                                                                                                                                                                                                                                                                                                                                           1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Clark H, Dennis K,
                                                                                                                                                                                             sequence of a thioredoxin polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 47; DB
100.0%; Pred. No. 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fong S,
                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Schoenfeld JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8;
                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 105;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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mucus; sputum; thioredoxin; active site; cohesive lung disease; cystic fibrosis; enzyme. Amino acid sequence of a thioredoxin polypeptide.

mucus; cohesive sputum;

Mus musculus. Synthetic.

25-MAR-2004

WO2004024868-A2

10-SEP-2003; 2003WO-US028526

밁 Ś

Conservative

0

Indels

٥,

Gaps

0

RESULT 66

ADM32935

ADM32935 standard; protein; 105

17-JUN-2004 ADM32935;

(first entry)

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The specification describes a method for increasing the liquefaction of CC mucus or sputum in a patient that has excessively viscous or cohesive CC mucus or sputum. The method comprises contacting the mucus or sputum of the patient with a composition comprises contacting the mucus or sputum of CC a thioredoxin active site in reduced state. The mucus or sputum to be CC contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the CC patient in a pharmaceutical carrier. The protein has a half-life in the CC patient of about 5 minutes to about 24 hours. The liquid phase of a total CC volume of a sample of mucus or sputum from the patient shows a CC statistically significant increase after administration of the CC composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin. The composition further comprises nicotinamide-adenine CC thioredoxin. The composition further comprises nicotinamide-adenine CC thioredoxin active site of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for CC excessive viscosity or cohesiveness of mucus or sputum is a symptom or CC cause of the disease. ADM32930ADM32941 represent thioredoxin polypeptides which may used in the method of the invention.
Query Match
Best Local S
Matches 6
                                                                                    Sequence 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 66-67; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     comprising a protein or peptide containing a thioredoxin active-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Increasing the liquefaction of excessively viscous or cohesive mucus sputum comprises contacting the mucus or sputum with a composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-270016/25.
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11-APR-2003; 2003US-0462082P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (NAJE-) NAT JEWISH MEDICAL & RES
  Similarity 6; Conser
                                                                                       AA;
100.0%; Score 47; DI
100.0%; Pred. No. 14.
cive 0; Mismatches
                         DB
14;
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                                            Length 105;
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RESULT 67
ADM32936
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the patient with a composition comprising a protein or peptide containing a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a total volume of a sample of mucus or sputum from the patient shows a statistically significant increase after administration of the composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian thioredoxin. The composition further comprises nicotinamide-adenine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       dinucleotide phosphate (reduced form) (NADPH) for reducing the thioredoxin active site of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for treating lung diseases (i.e. cystic fibrosis) in which abnormal or excessive viscosity or cohesiveness of mucus or sputum is a symptom or cause of the disease. ADM32930ADM32941 represent thioredoxin polypeptides which may used in the method of the invention.
                                                                                                                                                            mucus; sputum; thioredoxin; active site; cohesive mucus; cohesive lung disease; cystic fibrosis; enzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The specification describes a method for increasing the liquefaction of mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum. The method comprises contacting the mucus or sputum of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Increasing the liquefaction of excessively viscous or cohesive mucus or sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-SEP-2002; 2002US-0409960P.
11-APR-2003; 2003US-0462082P.
10-SEP-2003; 2003WO-US028526
                                                                                                                            Rattus norvegicus
                                                                                                                                                                                                                                                      17-JUN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                             ADM32936 standard; protein; 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 65; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-270016/25
                                    25-MAR-2004
                                                                        WO2004024868-A2
                                                                                                                                                                                                                  Amino acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NAJE-) NAT JEWISH MEDICAL & RES CENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6;
                                                                                                                                                                                                                                                                                                                                                                                                                         WCGPCK 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                  of a thioredoxin polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                               A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB
14;
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8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 105;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                 sputum;
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PR 11-APR-2002; 2002US-0462082P.

PR 11-APR-2003; 2003US-0462082P.

XX (NAJE-) NAT JEWISH MEDICAL & RES CENT.

XX White CW;

XI White CW;

XX WPI; 2004-270016/25.

XX WPI; 2004-270016/25.

XX WPI; 2004-270016/25.

XX Increasing the liquefaction of excessively viscous or cohesive mucus or sputum comprises contacting the mucus or sputum with a composition or comprising a protein or peptide containing a thioredoxin active-site in PT reduced state.

XX Disclosure; Page 65-66; 69pp; English.

XX The specification describes a method for increasing the liquefaction of mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum in a patient that has excessively rescue or cohesive contacted is located in the respiratory tract. The mucus or sputum of a thioredoxin active site in reduced state. The mucus or sputum of a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the patient in a pharmaceutical carrier me protein has a half-life in the patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a total composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thoredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thoredoxin selected from prokaryotic method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquifaction of excessively or camber of the protein and thioredoxin reductase. The composition are useful for decreasing the viscosity or cohesive mucus or sputum. These may be used for cause of the disease and any subdiving a symptom or cause of the disease and any subdiving a symptom or cause of the disease. Any 329 Janghyn 2241 regresent thioredoxin polypeptides which may used in the method of the invention.
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                á
                                         Query Match
Best Local Similarity
                                  Matches
31 WCGPCK 36
                1 WCGPCK 6
                                 6
                                 Conservative
                                          100.0%; Score 47; DB 100.0%; Pred. No. 14;
                                 <u>,</u>
                                  Mismatches
                                                  DB
                                                  8;
                                 0,
                                                 Length 105
                                 0,
                                Gaps
                                  0
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RESULT 68
ADM32937
ID ADM32937 standard; protein; 105 AA.

XX
AC ADM32937;
XX
DT 17-JUN-2004 (first entry)
XX
Mino acid sequence of a thioredoxin polypeptide.
XX
KW mucus; sputum; thioredoxin; active site; cohesive mucus; cohesive sputum;
KW lung disease; cystic fibrosis; enzyme.
XX
OS Bos taurus.
OS Synthetic.
XX
PN W02004024868-A2.
XX
PP 10-SEP-2003; 2003WO-US028526.
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ADM32934
ID ADM1
XX
AC ADM1
XX
DT 17-1
XX
DT 17-1
XX
DX Amii
XX
IU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mucus or sputum. The method comprises contacting the mucus or sputum of
the patient with a composition comprising a protein or peptide contraining
a thioredoxin active site in reduced state. The mucus or sputum to be
contacted is located in the respiratory tract, gastrointestinal tract or
reproductive tract of the patient. The composition is administered to the
patient of about 5 minutes to about 24 hours. The liquid phase of a total
composition. The protein comprises thioredoxin selected from prokaryotic
composition. The protein comprises after administeration of the
composition. The protein comprises thioredoxin selected from prokaryotic
thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian
of thioredoxin. The composition further comprises nicotinamide-adenine
chinucleotide phosphate (reduced form) (NADPH) for reducing the
thioredoxin active site of the protein, and thioredoxin reductase. The
method and composition are useful for decreasing the viscosity and
cohesiveness of, and increasing the liquideaction of excessively or
treating lung diseases (i.e. cystic fibrosis) in which abnormal or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      excessive viscosity or cohesiveness of mucus or sputum is a symptom or cause of the disease. ADM32930ADM32941 represent thioredoxin polypeptides which may used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Increasing the liquefaction of excessively viscous or cohesive mucus or sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 66; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-SEP-2002; 2002US-0409960P.
11-APR-2003; 2003US-0462082P.
   10-SEP-2003; 2003WO-US028526
                                                                   25-MAR-2004
                                                                                                                                 WO2004024868-A2
                                                                                                                                                                                                                                Gallus
                                                                                                                                                                                                                                                                                                                                                                                            Amino acid sequence of a thioredoxin polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADM32934 standard; protein; 105 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                                                                                                                                                                                                                                                             Lung
                                                                                                                                                                                                                                                                                                                                  mucus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                             disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Н
                                                                                                                                                                                                                            gallus
                                                                                                                                                                                                                                                                                                                           sputum; thioredoxin; active site; cohesive mucus; cohesive
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                                                                                                                                                                                                                                                                                          fibrosis; enzyme
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commous or sputum. The method comprises contacting the mucus or sputum. The method comprises contacting the mucus or sputum of the patient with a composition comprises contacting the mucus or sputum of the patient with a composition comprises contacting a protein or peptide containing a threadoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the patient of about 5 minutes to about 24 hours. The liquid phase of a total covolume of a sample of mucus or sputum from the patient shows a composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian composition further comprises nicotinamide-adenine thioredoxin active site of the protein, and thioredoxin and mammalian chioredoxin active site of the protein, and thioredoxin reductase. The composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquifaction of excessively or cohesive mucus or sputum. These may be used for treating lung diseases (i.e. cystic fibrosis) in which abnormal or cause of the disease. ADM32930ADM32941 represent thioredoxin polypeptides of which may used in the method of the invention.
Query Match
Best Local S
Matches 6
                                                                                       Sequence 105
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11-APR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The specification describes a method for increasing the liquefaction of mucus or sputum in a patient that has excessively viscous or cohesive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprising a protein or peptide containing a thioredoxin active-site
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ADO19655 RESULT 70 ADO19655 standard; protein; 105

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12-AUG-2004

(first entry)

Human; PRO; immune related disorder; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; juvennile chronic arthritis; systemic sclerosis; Sjogren's syndrome; vasculitis; sarcoidosis; autoimmune haemolyric anaemia; autoimmune thrombocytopenia; thyroiditis; diabetes mellitus; renal disease; demyelinating disease; central nervous system; peripheral nervous system; demyelinating polyneuropathy; Guillain-Barre syndrome; chronic inflammatory demyelinating polyneuropathy. Human PRO polypeptide #292.

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Search completed: February 23, 2006, 00:37:12 Job time : 163 secs
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                                                                                                                                               Query Match 100.0%; Score 47; DB 8; Length 105; Best Local Similarity 100.0%; Pred. No. 14; Matches 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                           The invention relates to human PRO polypeptides and the polynucleotides encoding them. The polypeptides and polynucleotides are useful for treating and diagnosing immune related disorders in mammals. The immune related disorders include systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis, systemic sclerosis, Sjogren's syndrome, vasculitis, sarcoidosis, autoimmune haemolytic anaemia, autoimmune thrombocytopenia, thyroiditis, diabetes mellitus, immune-mediated renal disease, demyelinating diseases of the central or peripheral nervous system, demyelinating polyneuropathy, Guillain Barre syndrome and chronic inflammatory demyelinating polypeptide of the inverse.
                                                                                                                                                                                                                            Sequence 105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel PRO polypeptide e.g., PRO69614, PRO71106, or PRO86388 useful for treating an immune related disorder such as systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis or spondyloarthropathy.
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N-PSDB; ADO19654.
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A; Title: The minimal gene complement of Mycoplasma genitalium. A; Reference number: A64200; MUID:96026346; PMID:7569993
A; Status: nucleic acid sequence not shown; translation not shown.
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R;Ng, W.V.; Kennedy, S.P.; Mahairas, G.G.; Berquist, B.; Pan, M.; Shukla, H.D.; Lasky, Leithauser, B.; Keller, K.; Cruz, R.; Danson, M.J.; Hough, D.W.; Maddocks, D.G.; JabJung, K.H.; Alam, M.; Freitas, T.

Proc. Natl. Acad. Sci. U.S.A. 97, 12176-12181, 2000
A;Authors: Hou, S.; Daniels, C.J.; Daniels, C.J.; Comer, A.D.; Ebhardt, H.; Lowe, T.M.; A;Title: Genome sequence of Halobacterium species NRC-1.
A;Reference number: A84160; MUID:20504483; PMID:11016950
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Best Local Similarity
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R;Hilmmelreich, R; Hilbert, H.; Plagens, H.; Pirkl, E.; Li, B.C.; Herrmann, R.
Nucleic Acids Res. 24, 4420-4449, 1996
A;Title: Complete sequence analysis of the genome of the bacterium Mycoplasma pneumoniae
A;Reference number: S73327; MUID:97105885; PMID:8948633
A;Accession: S73896
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Nolecule type: DNA
A;Residues: 1-102 <HIM>
A;Residues: 1-102 <HIM>
A;Cross-references: UNIFROT:P75512; UNIPARC:UPI0000136E8A; EMBL:AE000056; GB:U00089; NID
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, November 1996
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Residues: 1-102 <TIGR>
A;Cross-references: UNIPROT:P47370; UNIPARC:UPI0000136E89; GB:U39691; GB:L43967; NID:g10
A;Experimental source: strain G-37
C;Genetics:
A;Genetic code: SGC3
C;Superfamily: thioredoxin; thioredoxin homology
C;Kuperfamily: thioredoxin; thioredoxin homology
C;Kupords: redox-active disulfide
F;9-90/Domain: thioredoxin homology <THR>
F;9-90/Domain: thioredoxin homology <THR>
                                                                                                                         A;Status: preliminary
A;Molecule type: DNA
A;Molecule type: CARN
A;Residues: 1-10-4 ARN>
A;Residues: 1-10-4 ARN>
A;Cross-references: UNIPROT:084544; UNIPARC:UPI0000136E75; GB:AE001324; GB:AE001273; NII
A;Experimental source: serotype D, strain UW-3/Cx
C;Genetics:
                                                                                                                                                                                                                                                                                                                                          probable thioredoxin - Chlamydia trachomatis (serotype D, strain UW3/Cx) C;Species: Chlamydia trachomatis C;Species: Chlamydia trachomatis C;Date: 13-Sep-1998 #sequence_revision 13-Sep-1998 #text_change 05-Oct-2 C;Accession: B71503 R;Stephens, R.S.; Kalman, S.; Lammel, C.J.; Fan, J.; Marathe, R.; Aravin Science 282, 754-759, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Genetic code: SGC3
C;Superfamily: thior
F;9-90/Domain: thior
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                                                                                                                                                                                                                                                                      A;Title: Genome sequence of A;Reference number: A71570; A;Accession: B71503
                                                                           A; Gene: trxA
C; Superfamily:
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N;Alternate names: hypothetical protein A65_orf102
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                                                                                                                                                                                                                                                                                                     of an obligate intracellular pathogen 70; MUID:99000809; PMID:9784136
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A;Cross-references: UNIPARC:UPI0000171E21
R;Koetter, P.; Rose, M.; Entian, K.D.
submitted to the Protein Sequence Database,
                                                                                                                                                                                                                                                                                                                                       R;Muller, E.G.D.
J. Biol. Chem. 266, 9194-9202, 1991
A;Title: Thioredoxin deficiency in yeast prolongs S phase and A;Reference number: A39847; MUID:91225027; PMID:2026619
A;Accession: A39847
                                                                                                                                                          A;Reference number: $05793; MUID:72100583; PMID:4945270
A;Accession: $15360
                                                                                                                                                                                                                                   A;Cross-references: UNIPARC:UPI000004F943; R;Hall, D.E.; Baldesten, A.; Holmgren, A.; Eur. J. Biochem. 23, 328-335, 1971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Title: Yeast thioredoxin genes.
A;Reference number: A38669; MUID:91107668; PMID:1988444
A;Accession: S15048
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                  A; Reference number: S64863
A; Accession: S64870
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N;Alternate names: protein L1933; protein YLR043c
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A;Experimental source: strain Nigg (MoPn)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Reference number: A81500; MUID:20150255; PMID:10684935
A;Accession: C81660
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           thioredoxin
                                                                                                                        A; Residues: 26-34 < HAL>
                                                                                                                                               A; Molecule type: protein
                                                                                                                                                                                                                                                                                                         A;Residues: 1-103 <MUL>
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A; Residues: 1-103 <GAN>
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A; Residues: 1-102 <TET>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic Acids Res. 28, 1397-1406, 2000
                                                                                                                                                                                                                                                                                                                         A; Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;Species: Saccharomyces cerevisiae
;Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 05-Oct-2004
;Accession: S15048; A39847; S15360; S64870; B38669
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;Species: Chlamydia muridarum, Chlamydia trachomatis MoPn
;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 05-Oct-2004
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Salzberg,
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                                                                                          C; Date: 30-Juni-13086
C; Accession: A28086
A; Reference number: A; Accession: A28086
                  R;Johnson, R.S.; Mathews, W.R.; Biemann, K.; Hopper, S. J. Biol. Chem. 263, 9599-9597, 1988
A;Title: Amino acid sequence of thioredoxin isolated from A;Reference number: A28086; MUID:88257078; PMID:3164311
                                                                                                                          C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 30-Jun-1989 #sequence_revision 26-May-1994
                                                                                                                                                                                                                   RESULT 8
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                                                                                                                                                                          thioredoxin
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thioredoxin II - fission yeast (Schizosaccharomyces pombe)
N,Alternate names: thioredoxine 2
C;Species: Schizosaccharomyces pombe
C;Date: 03-Dec-1999 #sequence revision 03-Dec-1999 #text_change 09-Jul-2004
C;Accession: T39085; T46567
                                                                                               A; Map position: 1
C; Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                       A;Cross-references: UNIPARC:UPI000016205F; A;Experimental source: strain 972(-)
                                                                                                                                                                                                                                                                                                                                 A;Description: TRX2, a fission yeast stress protein. A;Reference number: Z23075
A;Accession: T46567
                                                                                                                                                                                                                                                                                                                                                                                                           A;Experimental source: strain 972h-; cosmid c7D4
R;Lenaers, G.; Perret, E.; Bonin, O.; Picard, A.; Caput,
submitted to the EMBL Data Library, October 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Reference number: Z21826
A; Accession: T39085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Gentles, S.; Churcher, C.M.; Wood, V.; Barrell, B.G.; Rajandream, M.A. submitted to the EMBL Data Library, September 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     F;9-90/Domain: thioredoxin homology <THR>
F;30-33/Disulfide bonds: redox-active #st
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A;Residues: 1-103 <KOE>
A;Cross-references: UNIPARC:UPI000004F943; EMBL:Z73215; NID:g1360372; PIDN:CAA97572.1; |
A;Note: experimental_source strain S288C
C;Genetics:
                                                                                                                                                       A;Gene: trx2; SPAC7D4.07c
                                                                                                                                                                                                                                                                                                         A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: UNIPROT:014463; UNIPARC:UPI000016205F; EMBL:Z99532; PIDN:CAB16724.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Status: preliminary; translated from GB/EMBL/DDB
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  Similarity 6; Conserv
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                             Score 47;
Pred. No.
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Pred. No. 1;
Mismatches
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#text_change 05-Oct-2004

rabbit bone

marrow determined

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A; ROLECULE CYPE: DANA
A; CROSS-references: UNIPROT: P14949; UNIPARC: UPI00006087B; GB: J03294; GB: M26384; NID: G1
R; Kunst, F; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Berter
C.; Bron, S.; Browillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Chd
A.; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E.
A.; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E.
Nature 390, 249-256, 1997
A; Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Galler
iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hullo, M.F.
Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois,
A; Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maueel
y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maueel
y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.M.; Levine, A.; Ton, Portetelle
Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon,
A; Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Serot
akeuchi, M.; Tamaka, T.; Tranaka, T.; Trepstra, P.; Toganoi, A.; Tosato, V.; Uchiyama,
T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, K
A; Authors: Yoshikawa, H.F.; Zumstein, E.; Yoshikawa, H.; Danchin, A.
A; Title: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis.
A; Accession: H69726
A. Steries not selvence of the Gram-positive bacterium Bacillus subtilis.
                                                                                                                                                                                                                                                                                                                                                                                               A; Map position: 70 min
C; Superfamily: thioredox
C; Keypords: redox-active
F; 9-90/Domain: thioredox
F; 29-32/Disulfide bonds:
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C;Date: 31-Jan-1992 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
C;Accession: B37192; H69726
R;Chen, N:V; Zhang, J.J.; Paulus, H.
J. Gen. Microbiol. 135, 2931-2940, 1989
A;Title: Chromosomal location of the Bacillus subtilis aspartokinase II gene A;Reference number: A37192; MUID:90132525; PMID:2559145
A;Accession: B37192.
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thioredoxin
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                                                           RESULT 10
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Best Local
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                               29-32/Disulfide bonds:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Molecule type: DNA
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in: thioredoxin homology <THR>
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yeast
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(Saccharomyces
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Pred. No.
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Pred. No. 1;
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   cerevisiae)
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   A; Map position:
C; Superfamily: t
C; Keywords: redo
                                                                                       A;Gene: SGD:TRX2; TR-I; MIPS:YGR209c
A;Cross-references: SGD:S0003441; MIPS:YGR209c
   ;Superfamily: thioredoxin; thioredoxin; Keywords: redox-active disulfide
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A;Reference number: S53922
A;Accession: S53932
A;Molecule type: DNA
A;Residues: 1-104 <GUE>
A;Cross-references: UNIPARC:UPI000004F91C; EMBL:Z49133; NID:g790489; PIDN:CAA89002.1;
A;Cross-references: UNIPARC:UPI000004F91C; EMBL:Z49133; NID:g790489; PIDN:CAA89002.1;
A;Experimental source: strain S288C
R;Song, J.M.; Cheung, E; Rabinowitz, J.C.
submitted to the EMBL Data Library, November 1995
A;Description: Analysis of the 15.6-kb fragment encompassing the ADE3 gene.
A;Reference number: S61947
A;Rocession: S61947
A;Accession: S61947
A;Accession
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A,Description: Sequencing of a 17.6 kb segment on the right arm of yeast chromosome VII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPARC:UPI000004F91C; GB:M62648; NID:9
R;Hall, D.E.; Baldesten, A.; Holmgren, A.; Reichard, P.
Eur. J. Biochem. 23, 328-335, 1971
A;Title: Yeast thioredoxin. Amino-acid sequence around the A;Reference number: S05793; MUID:72100583; PMID:4945270
A;Accession: S05793
A;Molecule type: protein
A;Residues: 2;27-43;88-104 -HALD
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A;Title: Yeast thioredoxin genes.
A;Reference number: A38669; MUID:91107668;
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C;Species: Saccharomyces cerevisiae
C;Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 05-Oct-2004
C;Accession: S15049; B39847; S05793; S53932; S61947; S64531; S63858; A38669
                                                                                                                                                                                                                                                                                         Yeast 12, 273-280, 1996
Yeast 12, 273-280, 1996
A; Title: Sequencing of a 17.6 kb segment on the right arm of yeast terial electron-transferring flavoproteins (beta-chain) and of the terial electron-transferring 180000019; PMID:8904340
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: Thioredoxin deficiency in yeast prolongs S phase and A;Reference number: A39847; MUID:91225027; PMID:2026619 A;Accession: B39847
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A; Residues: 1-104 < GUW>
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A; Accession: S64531
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A; Residues: 1-104 < GAN>
     A;Cross-references: UNI
A;Note: the nucleotide
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A; Residues: 1-104 < GUF>
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                                                         UNIPARC: UPI000004F91C; EMBL: Z49133; NID: g790489; PIDN: CAA89002.1;
sequence was
     submitted to the EMBL Data Library, April 1995
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A;Cross-references: UNIPARC:UPI0000174DEF
R;Hammel, K.E.; Cornwell, K.L.; Buchanan, B.B.
Proc. Natl. Acad. Sci. USA 80, 3681-3685, 1983
A;Title: Ferredoxin/flavoprotein-linked pathway for the reduction of thioredoxin.
                                                                                                                                                           A;Cross-references: UNIPARC:UPI0000174DEF
R;Reynolds, C.M.; Poole, L.B.; Hammel, K.E.;
submitted to the Protein Sequence Database, S
A;Reference number: A59394
A;Accession: A59394
                                                                                                                                                                                                                                                                                                                                                                                                       thioredoxin - Clostridium pasteurianum C;Species: Clostridium pasteurianum C;Species: Clostridium pasteurianum C;Date: 02-Apr-2002 #sequence_revision 02-Apr-2002 #text_change 05-Oct-2004 C;Paccession: B59394; A59394 R;Reynolds, C.M.; Meyer, J.; Poole, L.B. Biochemistry 41, 1990-2001, 2002
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Mol. Microbiol. 16, 955-967, 1995
A;Title: Exploring the Mycoplasma capricolum genome: a minimal
A;Title: Exploring the Mycoplasma capricolum genome: a minimal
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F;31-34/Disulfide bonds: redox-active #status experimental
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                                                                                            A; Molecule type: protein A; Residues: 1-104 < RE2>
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A;Accession: B59394
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Reference number: S77739; MUID:96059641; A;Accession: S77780
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                                                                                                                                        A;Status: preliminary
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-13-93/Domain: thioredoxin homology <THR>
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Bork, P.; Ouzounis, C.; C
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R;Kuroda, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.; Cui, L.; ma, A.; Mizutani-Ui, Y.; Kobayashi, N.; Sawano, T.; Inoue, R.; Kaito, C.; Sekimizu, C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K.
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                                                                                                                                                                              C; Superfamily: thioredoxin; thioredoxin homology
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                                                                                                                                                                                                                                                 A;Cross-references: UNIPROT:Q9ZEH4; UNIPARC:UPI00000D78B6; GB:BA000018; PID:g13700945; A;Experimental source: strain N315
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A; Residues: 1-104 < KUR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                 C.; Shiba, T.; Hattori, M.
Lancet 357, 1225-1240, 2001
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C;Species: Bacillus halodurans
C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 05-Oct-2004
C;Accession: B84037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B84037
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               thioredoxin [imported] - Staphylococcus aureus (strain N315)
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                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;Cross-references: UNIPROT:Q9KBA8; UNIPARC:UPI00000C40CD; GB:AP001517; GB:BA000004; NII
;Experimental source: strain C-125
                                                                                                                                   Query Match
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28
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Pred. No.
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Pred. No.
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A; Martin, H.; Dean, M. Biochem. Biochem. Biochem. Sophys. Res. Commun. 175, 123-128, 1991
A; Title: Identification of a thioredoxin-related protein associated with plasma A; Reference number: PT0079; MUID:91151337; PMID:1998498
A; Accession: PT0079
                                                                                                 R;Jones, S.W.; Luk, K.C.
J. Biol. Chem. 263, 9607-9611, 1988
A;Title: Isolation of a chicken thioredoxin cDNA clone:
A;Reference number: A30006; MUID:88257080; PMID:2838473
A;Accession: A30006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: UNIPROT:P08629; UNIPARC:UPI00001713BC; C;Superfamily: thioredoxin; thioredoxin homology C;Keywords: redox-active disulfide
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C;Date: 31-Mar-1989 #sequ
                                                                                                                                                                                                                                                                   A;Cross-references: UNIPARC:UPI000011065C; EMBL:X77584; NID:g453963; PIDN:CAA54687.1; PI R;WOllman, E.E.; d'Auriol, L.; Rimsky, L.; Shaw, A.; Jacquot, J.P.; Wingfield, P.; Grabe J. Biol. Chem. 263, 15506-15512, 1988
A;Title: Cloning and expression of a cDNA for human thioredoxin.
A;Reference number: A31993; MUID:89008454; PMID:3170595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gene 102, 221-228, 1991
A;Title: Isolation and characterization of human thioredoxin-encoding genes A;Reference number: JH0568; MUID:91340156; PMID:1874447
A;Accession: JH0568
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Species: Homo sapiens (man)
C;Date: 30-Jun-1992 #sequence revision 26-May-1994 #text change 05-Oct-2004
C;Accession: JH0568; S04106; S44375; A31993; PT0079; A60749; A38922; S53453
R;Tonissen, K.F.; Wells, J.R.E.
Gene 102, 221-228, 1991
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A; Residues: 1-105 < JON>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: UNIPROT:P10599; UNIPARC:UPI000011065C; EMBL:X54539; NID:g37455; PIDN R;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kondo, N.; Matsui, H.; Hamuro, J.; Brown, N.; Arai EMBO J. 8, 757-764, 1989
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A; Residues: 1-105 < TAG2 >
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Contents: erratum A; Accession: S44375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: UNIPARC:UPI000011065C; GB:X77584; NID:g453963; PIDN:CAA54687.1; PID A;NOte: this sequence has been revised in reference S44375 R;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kando, N.; Matsui, H.; Hamuro, J.; Brown, N.; Arai EMBO J. 13, 2244, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Title: ATL-derived factor (ADF), an IL-2 receptor/Tac A;Reference number: S04106; MUID:89251607; PMID:2785919 A;Accession: S04106
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A;Residues: 1-105 <TON>
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100.0%; Pred. No.
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                                                                                                                                                        GB:J04026; NID:g339648; PIDN:AAA74596.1;
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A; Molecule type: protein

A; Residues: 1-21;38-57 < DBA>

A; Residues: 1-21;38-57 < DBA>

A; Residues: 1-21;38-57 < DBA>

A; Cross-references: UNIPARC: UPI00000353C5; UNIPARC: UPI0000171B1C

A; Note: described to be a surface-associated thioredoxin

R; Wakasugi, H.; Rimsky, L.; Mahe, Y.; Kamel, A.M.; Fradelizi, D.; Tu

Proc. Natl. Acad. Sci. U.S.A. 84, 804-808, 1987

A; Title: Epstein-Barr virus-containing B-cell line produces an inter

A; Reference number: A60870; MUID: 87118252; PMID: 3027706

A; Reference number: A60870; MUID: 87118252; PMID: 3027706

A; Contents: annotation

R; Weichsel, A.; Gasdaska, J.R.; Powis, G.; Montfort, W.R.

submitted to the Brookhaven Protein Data Bank, February 1996

A; Reference number: A6553; PDB: IBRT

A; Contents: annotation; X-ray crystallography, 1.7 angstroms, reduce

R; Weichsel, A.; Gasdaska, J.R.; Powis, G.; Montfort, W.R.

Schwisted to the Brookhaven Protein Data Bank, February, 1006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    submitted to the Brookhaven Protein Data Bank, February 1996
A;Reference number: A65534; PDB:1ERU
A;Contents: annotation; X-ray crystallography, 2.1 angstroms, oxidized forn
R;Forman-Kay, J.D.; Clore, G.M.; Gronenborn, A.M.
submitted to the Brookhaven Protein Data Bank, December 1990
A;Reference number: A50924; PDB:4TRX
A;Rontents: annotation; conformation by (1)H-NMR, residues 1-73,'T',75-105
R;Forman-Kay, J.D.; Clore, G.M.; Wingfield, P.T.; Gronenborn, A.M.
Biochemistry 30, 2685-2698, 1991
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R;Dean, M.F.; Martin, H.; Sansom, P.A.
Biochem. J. 304, 861-867, 1994
A;Title: Characterization of a thioredoxin-related surface protein.
A;Title: Characterization of a thioredoxin-related surface protein.
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A; Residues: 2-12, 'K', 14-15, 'XX', 18-19, 'X', 21-22 <SIL>
A; Cross-references: UNIPARC: UPI0000171E1A
A; Cross-references: UNIPARC: UPI0000171E1A
A; Note: the abstract is inconsistent with figure 4 in having one undetermined residue
B; Rimsky, L.; Wakasugi, H.; Ferrara, P.; Robin, P.; Capdevielle, J.; Tursz, T.; Fradel
J. Immunol. 136, 3304-3310, 1986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: UNIPARC:UPI0000171E19
R;Silberstein, D.S.; Ali, M.H.; Baker, S.L.; David, J.R.
J. Immunol. 143, 979-983, 198
A;Title: Human eosinophil cytotoxicity-enhancing factor.
A;Reference number: A60749; MUID:89309777; PMID:2745979
A;Accession: A60749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: High-resolution three-dimensional structure of reduced recombinant A;Reference number: A38953; MUID:91159399; PMID:2001356 A;Contents: annotation; conformation by (1)H- and (15)N-NMR C;Comment: This small ubiquitous protein functions in many intracellular bic
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                                                                                                                                                                                                                                                                                                                                                                       A;Map position: 9q31-9q31
A;Introns: 8/3; 43/3; 63/3; 85/3
C;Superfamily: thioredoxin; thioredoxin homology
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A; Residues: 2-16 < WAK>
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                                                                                                                                                                                                                                                                                               C;Keywords: redox-active disulfide F;2-105/Product: thioredoxin #status experimental
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                                                                                                                                                                                                                          F;32-35/Disulfide bonds:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: GDB:120475; OMIM:187700
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                                                                                Matches
                                                                                                                  Best Local
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   1 WCGPCK 6
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C;Superfamily: thioredoxin; thioredoxin homology C;Keywords: redox-active disulfide F;9-92/Domain: thioredoxin homology <THR> F;32-35/Disulfide bonds: redox-active #status predicted
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A;Molecule type: mRNA
A;Residues: 1-93,'N',94-96,'ALT',100-104,'S' <TAG2>
A;Residues: UNIPARC:UPI0000171E1E; GB:X77585
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A;Title: Thioredoxin gene expression is transcriptionally up-regulated A;Reference number: J80667; MUID:92181438; PMID:1543487
A;Accession: J80667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Title: ATL-derived factor (ADF), an IL-2 receptor/Tac A;Reference number: S04106; MUID:89251607; PMID:2785919 A;Accession: S04107
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A;Title: Structure of the mouse thioredoxin-encoding gene and its A;Reference number: JC4068; MUID:95137382; PMID:7835695
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C;Date: 21-Nov-1993 #sequence_revision 17-Oct-1997 #text_change
C;Accession: JC4068; S44376; S04107
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N;Alternate names: ATL-derived factor (ADF)
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A; Residues: 1-105 < ANG>
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C;Species: Macaca mulatta (rhesus macaque)
C;Date: 30-Jun-1992 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
C;Accession: JS0667
                                                                                                                                                                                                                                                                                                                                                      A; Map position: 4:24.6
A; Introns: 29/2; 44/1; 84/2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: mRNA
A; Residues: 1-105 < TAG1>
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A; Residues: 1-105 < MAT>
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                             100.0%; Score 47; llarity 100.0%; Pred. No. Conservative 0; Mismatch
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100.0%; Pred. No. 1;
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                             Gaps
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RESULT 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             thioredoxin - coryneform bacterium ATCC11425
c;Specias: coryneform bacterium ATCC11425
C;Date: 02-Apr-1982 #sequence_revision 02-Apr-1982 #text_change 05-Oct-2004
C;Accession: A00281
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Biochem. J. 304, 861-867, 1994
A;Title: Characterization of a thioredoxin-related surface protein.
A;Reference number: S53453; MUID:95118305; PMID:7818492
                                                                                                                                                  C; Keywords: redox-active disulfide F;8-91/Domain: thioredoxin homology <THR>F;30-33/Disulfide bonds: redox-active #st:
                                                                                                                                                                                                                                                A;Cross-references: UNIPROT:P00275; UNIPARC:UPI0000136D69
A;Note: the source was designated as Corynebacterium nephridii
C;Comment: Thioredoxins are ubiquitous small hydrogen carrier
                                                                                                                                                                                                                                                                                                                                A; Molecule type: protein A; Residues: 1-105 < MEN>
                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Meng, M.; Hogenkamp, H.P.C.
J. Biol. Chem. 256, 9174-9182, 1981
A;Title: Purification, characteriza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                F;32-35/Disulfide bonds: redox-active #status predicted
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A; Residues: 1-105 < TON>
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                                                                                                                                                                                                                  Comment: Thioredoxins are uniquitous swart ayur Superfamily: thioredoxin thioredoxin homology
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100.0%; Pred. No.
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Pred. No. 1;
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29

WCGPCK

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thioredoxin
C;Species: St
C;Date: 06-Ja
C;Accession:
R;Labudova, (
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A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-105 < KUR>
A;Cross-references: UNIPROT:Q97EM7; UNIPARC:UPI00000CA715; GB:AE001437; PIDN:AAK81023.1
A;Cross-references: UNIPROT:Q97EM7; UNIPARC:UPI00000CA715; GB:AE001437; PIDN:AAK81023.1
A;Cross-references: UNIPROT:Q97EM7; UNIPARC:UPI00000CA715; GB:AE001437; PIDN:AAK81023.1
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A;Title: Mechanisms of Evolution in Rickettsia conorii and Rickettsia prowazekii.
A;Reference number: A97700; MUID:21442074; PMID:11557893
A;Accession: B97700
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C;Species: Rickettsia conorii
C;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text chance.
                                                                                                                                                                                                                 RESULT 23
S33357
A;Description: PCR cloning
A;Reference number: $33357
A;Accession: $33357
A;Molecule type: DNA
A;Residues: 1-106 <LAB>
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C;Superfamily: thioredoxin; thioredoxin homology
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A;Molecule type: DNA
A;Residues: 1-105 <KUR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .; Daly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D. Bacteriol. 183, 4823-4838, 2001
                                                                                                nioredoxin - Streptomyces aureofaciens (fragment)
;Species: Streptomyces aureofaciens
;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change
;Accession: S33357
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A;Cross-references: UNIPROT:P56430; UNIPARC:UPI0000136E82; GB:AE000594; GB:AE000511; NIT A;Experimental source: strain 26695
R;Alm, R.A.; Ling, L.S.L.; Moir, D.T.; King, B.L.; Brown, E.D.; Doig, P.C.; Smith, D.R., I Ives, C.; Gibson, R.; Merberg, D.; Mills, S.D.; Jiang, Q.; Taylor, D.E.; Vovis, G.F.; Nature 397, 176-180, 1999
A;Title: Genomic sequence comparison of two unrelated isolates of the human gastric path A;Reference number: A71800; MUID:99120557; PMID:9923682
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A;Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser, A;Title: The complete genome sequence of the gastric pathogen Helicobacter pylori. A;Reference number: A64520; MUID:97394467; PMID:9252185
A;Accession: H64622
                                                                                                                                                                                                    R;Cohen, G.; Argaman, A.; Schreiber, R.; Mislovati, M.; Aharonowitz, J. Bacteriol. 176, 973-984, 1994
A;Title: The thioredoxin system of Penicillium chrysogenum and its po
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C; Superfamily: thioredoxin; thioredoxin homology
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A; Residues: 1-106 < ARN>
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A; Residues: 1-106 < TOM>
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A; Molecule type: DNA
A; Residues: 1-106 < COH>
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C;Species: Penicillium chrysogenum
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A;Variety: strains J99, 26695
                                                                            A;Status: preliminary
                                                                                                                   A; Accession: A49888
                                                                                                                                                         A; Reference number:
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A;Cross-references:

UNIPROT: P34723; UNIPARC: UPI0000136E91;

EMBL: X76120; NID: g426468;

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R;Wood, D
erage, G.
                                                                                                     A; Molecule type: protein
A; Residues: 1-107 < JOH>
A; Residues: 1-107 < JOH>
A; Cross-references: UNIPARC: UPI0000136E76
A; Cross-references: UNIPARC: UPI0000136E76
A; Note: unidentified residues are Ile or Leu
C; Superfamily: thioredoxin; thioredoxin homology
C; Keywords: heat-stable protein; redox-active disulfide
C; Reywords: heat-stable protein; redox-active disulfide
F; 10-93/Domain: thioredoxin homology < THR>
                                                                                                                                                                                                                                                                        Biochemistry 26, 1209-1214, 1987
A; Title: The primary structure of thioredoxin from Chromatium A; Reference number: A26622; MUID:87185419; PMID:3567166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cnioredoxin C-1 trxA [imported] - Agrobacterium tumefaciens (strain C58, Du
C;Species: Agrobacterium tumefaciens
C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 05-Oct-2004
C;Accession: AG2579
                                                                                                                                                                                                                                                                                                                                                              C;Species: Chromatium vinosum
C;Date: 31-Mar-1988 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
C;Accession: A26622
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Authors: Yoo, H.; Tao, ster, E.W.
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C;Superfamily: thioredoxin;
F;10-92/Domain: thioredoxin
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Science 294, 2317-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Gene: trxA
                                                                                                                                                                                                                                                                                                                                        R; Johnson, R.S.; Biemann, K.
                                                                                                                                                                                                                                                                                                                                                                                                                            thioredoxin
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A;Experimental source: strain C58 (Dupont)
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                                                                                        32-35/Disulfide
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    100.0%; Score 47; DB 1; Length 107; llarity 100.0%; Pred. No. 1.1; Conservative 0; Mismatches 0; Indels
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                                                                                 redox-active #status predicted
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Pred. No. 1.1;
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Mismatches
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M.; McClel
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thioredoxin C-1 [imported] - Brucella melitensis (strain 16M)
C;Species: Brucella melitensis
C;Date: 01-Feb-2002 #sequence_revision 01-Feb-2002 #text_change 05-Oct-2004
C;Accession: AH3504
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A;Title: The Drosophila maternal effect locus deadhead encodes a thioredoxin homolog reconstruction of the property of the prop
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C; Superfamily: thioredoxin; thioredoxin homology
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A;Residues: Î-107 <PYL>
A;Cross-references: UNIPROT:Q25345; UNIPARC:UPI000007DADA; GB:AE001274; NID:g3264850; PI
A:Experimental source: strain MHOM/IL/81/Friedlin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Reference number: A81455; MUID:99178987; PMID:10077609
A;Accession: B81458
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A;Title: Leishmania major Friedlin chromosome 1 has an unusual distribution of protein-c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Date: 24-Mar-1999 #sequence_revision 24-Mar-1999 #text_change 09-Jul-2004 C;Accession: B81458; T02814
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C;Keywords: redox-active disulfide
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C;Species: Drosophila melanogaster
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change
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                                                                                                                                                                    AH3504
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C;Species: 1
C;Date: 18-1
C;Accession
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A;Authors: Gnehm, C.L.; McDonald, L.A.; Small, K.V.; Fraser, C.M.; Smith, H.O.; Venter, A;Title: Whole-genome random sequencing and assembly of Haemophilus influenzae Rd. A;Reference number: A64000; MUID:95350630; PMID:7542800
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E64047
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A;Title: The genome sequence of the facultative intracellular pathogen Brucella melitens A;Reference number: AD3252; PMID:11756688
A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic pathon A;Reference number: A82950; MUID:20437337; PMID:10984043
A;Accession: G82991
A;Status: preliminary
A;Molecule type: DNA
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F;32-35/Disulfide bonds: redox-active #status predicted
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C;Species: Haemophilus influenzae
C;Date: 18-Aug-1995 #sequence_revision 18-Aug-1995 #text_change 05-Oct-2004
C;Accession: E64047
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C; Superfamily:
                                                                                                                             R;Stover, C.K.; Pham, X.Q.; Erwin, adman, S.; Yuan, Y.; Brody, L.L.; C.; Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
                                                                                                                                                                                                                                    thioredoxin pA5240 [imported] - Pseudomonas aeruginosa (strain PAO1) C;Species: Pseudomonas aeruginosa (c;Species: Pseudomonas aeruginosa (c;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-JC;Accession: G82991
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A; Residues: 1-107 < KUR>
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A; Residues: 1-107 < TIGR >
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Best Local
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K.R.; Kas, A.;
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A;Residues: 1-108 <STO>
A;Cross-references: UNIPROT:Q9X2T1; UNIPARC:UPI0000136E95; GB:AE004936; GB:AE004091; NID A;Experimental source: strain PAO1
A;Experimental source: crain PAO1
C;Genetics:
A;Gene: trxA; PA5240
C;Superfamily: thioredoxin; thioredoxin homology
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R;Heidelberg, J.F.; Eisen, J.A.; Nelson, W.C.; Clayton, R.A.; Gwinn, M.L.; Dodson, R.J. chardson, D.; Ermolaeva, M.D.; Vamathevan, J.; Bass, S.; Qin, H.; Dragoi, I.; Sellers, l, R.R.; Mekalanos, J.J.; Venter, J.C.; Fraser, C.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   thioredoxin VC0306 [imported] - Vibrio cholerae (strain N16961 serogroup O1) C;Species: Vibrio cholerae (c;Species: Vibrio cholerae (c;Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 05-Oct-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; Prentice, deno-Tarraga, A.M.; Chillingworth, T.; Cronin, A.; Davies, R.M.; Davis, P.; Dougan, il, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; Whitehead, S.; Barre
                                                                                                                                                                                                                           A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-108 <HEI>
                                                                                                                                                                                                                                                                                                                         Nature 406, 477-483, 2000
A;Title: DNA Sequence of both chromosomes of the cholera pathogen Vibrio cholerae.
A;Reference number: A82035; MUID:20406833; PMID:10952301
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C;Genetics:
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                                                                                                                                                                                  A;Cross-references: UNIPROT:Q9KV51; UNIPARC:UPI00000C2C6E; GB:AE004119; GB:AE003852; A;Experimental source: serogroup O1; strain N16961; biotype El Tor
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                     100.0%; Score 47; DB 100.0%; Pred. No. 1.1;
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R;Miloso, M.; Limauro, D.; Alifano, P.; Rivellini, F.; Lavitola, A.; Gulletta, E.; Brun: J. Bacteriol. 175, 8030-8037, 1993
A;Title: Characterization of the rho genes of Neisseria gonorrhoeae and Salmonella typh: A;Reference number: A49917; MUID:94075245; PMID:8253691
A;Accession: A49917
              A;Molecule type: DNA
A;Residues: 'MLHQQRNQHARLIPVELY',1-109 <BLAT>
A;Cross-references: UNIPARC:UPI0000165538; GB:AE000454; GB:U00096; NID:g2367278;
                                                                                                                                                                                                                                                                                                                                               A;Title: Genetic and physical analysis of the thioredoxin (trxA) A;Reference number: A91519; MUID:85155506; PMID:6099324 A;Accession: A91519
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A;Note: the nucleotide sequence was submitted to the EMBL Data Library, March 1992
R;Miloso, M.; Limauro, D.; Alifano, P.; Rivellini, F.; Lavitola, A.; Gulletta, E.; Bruni J. Bacteriol. 175, 8030-8037, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Kotani, H.; Nakajima, K.
Nucleic Acids Res. 20, 1424, 1992
A;Title: Cloning and sequence of thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 35
S35497
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Date: 24-Apr-1984 #sequence_revision 01-Sep-1995 #text_change 05-Oct-2004 
C;Accession: A91519; H65181; S30676; I54863; A91802; I52550; A91236; A00280; A22425;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Reference number: S35497; MUID:92220625; A;Accession: S35497
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Residues:
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                                                                                     A;Status: nucleic acid sequence not shown; translation not shown
                                                                                                              A; Accession: H65181
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;Date: 09-Dec-1993 #sequence revision;
;Accession: S35497; A49917; S31928
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     [validated] - Escherichia coli (strain K-12)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1-109 <MIL>
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source: strain
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Pred. No. 1.1;
0; Mismatches
substrain MG1655
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                                                               A; Molecule type: DNA
A; Residues: 1-109 < L
A; Experimental source: strain K12
                                    A;Cross-references: UNIPARC:UPI000003112A; GB:M10424; NID:g147608; PIDN:AAA24533.1;
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A; Residues: 'MLHQQRN
                                                                                                                                                                                                                                                                                                                                                                                   A;Status: translation not shown; translated from GB/EMBL/DDBJ
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A;Note: the sequence represents translation from an upstream ATG triplet that seems R;Daniels, D.L.; Plunkett III, G.; Burland, V.; Blattner, F.R. Science 257, 771-778, 1992
A;Title: Analysis of the Escherichia coli genome: DNA sequence of the region from 84
A;Reference number: S30660; MUID:92358234; PMID:1379743 not

A; Accession: S30676

A; Note: the sequence represents translation from an upstream R; Matsumoto, Y; Shigesada, K.; Hirano, M.; Imai, M. PROISECULE TYPE: THE CONTROL OF TRANSPORT OF THE PROPERTY (1-109 CDANS)

CROSS-references: UNIPARC:UPI0000165538; EMBL:M87049; NID:9836656; PIDN:AAA67582.1; PI

CROSS-references: UNIPARC:UPI0000165538; EMBL:M87049; NID:9836656; PIDN:AAA67582.1; PI

CROSS-references: UNIPARC:UPI0000165538; EMBL:M87049; NID:9836656; PIDN:AAA67582.1; PI

Bacteriol. 166, 945-958, 1986

A;Title: Autogenous regulation of the gene for transcription termination factor rho in A;Reference number: I54863; MUID:86223816; PMID:2423505 A;Accession: I54863

A; Molecule type: DNA A; Residues: 1-109 < M 1-109 <MAS>

A;Cross-references: UNIPARC:UPI000003112A; R;Lim, C.J.; Geraghty, D.; Fuchs, J.A. J. Bacteriol. 163, 311-16, 1985 GB:M12779; NID:g148067; PIDN:AAA24694.1; PID

PMID:3891733 о Е Escherichia

A;Title: Cloning and nucleotide sequence of the trxA A;Reference number: A91802; MUID:85234377; PMID:3891; A;Accession: A91802

1-109 <LIM>

PID

Escherichia coli

Molecule type: DNA

A;Residues: A;Cross-references: UNIPARC:UPI000003112A; GB:M26133; NID:g148065; PIDN:AAA24693.1; 1-109 <HOE>

PID

R;Holmgren,

A; Reference number: A91236; A; Title: Thioredoxin. . Biochem. 6, 475-484, 1968 . Biochem. 6. The amino acid sequence of the protein e: Thioredoxin. 6. The amino acid sequence of the protein vence number: A91236; MUID:69079993; PMID:4883076 from Escherichia coli

A; Molecule type: protein
A; Residues: 2-16, 'IV', 19-71, 'IG', 74-109 <HOL>
A; Cross-references: UNIPARC: UPI0000171E22 Accession: A91236

source: strain

A; Experimental

R;Haeberlein, eberiein, i. J. Biochem. 223, 473-479, 1994

A;Title: Structure requirements for disulfide bridge sulfitolysis of oxidized Escherichi A;Reference number: \$45671; MUID:9433336; PMID:8055916
A;Contents: annotation; chemical activity of wild type and engineered sequence R;Katti, S.K.; LeMaster, D.M.; Eklund, H.
submitted to the Brookhaven Protein Data Bank, March 1990
A;Reference number: A50900; PDB:2TRX
A;Contents: annotation; X-ray crystallography, 1.68 angstroms, residues 2-109
A;Katti, S.K.; LeMaster, D.M.; Eklund, H.
J. Mol. Biol. 212, 167-184, 1990
A;Title: Crystal structure of thioredoxin from Escherichia coli at 1.68 Angstroms resolu

A; Reference number: A58630; MUID: 90204538; PMID: 2181145

A;Contents: annotation; X-ray crystallography, 1.68 angstroms R;Holmgren, A.; Soderberg, B.O.; Eklund, H.; Branden, C.I. Proc. Natl. Acad. Sci. U.S.A. 72, 2305-2309, 1975

ö 2.8

angstrom

A;Title: Three-dimensional structure of Escherichia coli thioredoxin-S-2 A;Reference number: A93800; MUID:75176930; PMID:1094461 A;Contents: annotation; X-ray crystallography, 2.8 angstroms R;Dyson, H.J.; Gippert, G.P.; Case, D.A.; Holmgren, A.; Wright, P.E. submitted to the Brookhaven Protein Data Bank, January 1990 A;Reference number: A50871; PDB:1TRX

A;Contents: annotation; conformation by (1)H-NMR, reduced form, R;Dyson, H.J.; Gippert, G.P.; Case, D.A.; Holmgren, A.; Wright, residues P.E. 2-109

BIOCHEMISTRY 29, 4129-4136, 1990
A;Title: Three-dimensional solution structure of the reduced A;Reference number: A58631; MUID:90298180. form of Escherichia coli ţ,

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C.Accession: S27053
C.Accession: S27053
C.Accession: S27053
R;le Marechal, P.; Hoang, B.M.C.; Schmitter, J.M.; van Dorsselaer, A.; Decottignies, Ryr. J. Biochem. 210, 421-429, 1992
A;Title: Purification, properties and primary structure of thioredoxin from Aspergil. A;Reference number: S27053; MUID:93092975; PMID:1459127
             R; Parkhill, J.; Dougan, G.; James, K.D.; Thomson, N.R.; Pickard th, T.; Connerton, P.; Cronin, A.; Davis, P.; Davies, R.M.; Dow S.; Moule, S.; O'Gaora, P.
Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quall, M.; Rutherford, K.; Simmonds, M.; A;Title: Complete genome sequence of a multiple drug resistant A;Reference number: AB0502; MUID:21534947; PMID:11677608
                                                                                                                                                              thioredoxin [imported] - Salmonella enterica subsp. enterica serovar Typhi C;Species: Salmonella enterica subsp. enterica serovar Typhi A;Note: this species has also been called Salmonella typhi C;Date: 09-Nov-2001 #sequence_revision 09-Nov-2001 #text_change 05-Oct-2004 C;Accession: AF0922
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Contents: annotation; conformation by (1)H-, (15)N-NMR, oxidized form, residues R;Jeng, M.F.; Campbell, A.P.; Begley, T.; Holmgren, A.; Case, D.A.; Wright, P.E.; submitted to the Brookhaven Protein Data Bank, November 1995
A;Reference number: A66948; PDB:1XOB
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R;Jeng, M.F; Campbell, A.F; Begley, T.; Holmgren, A.; Case, D.A.; Wright, P.E.;
submitted to the Brookhaven Protein Data Bank, November 1995
A;Reference number: A66947; PDB:1XOA
A; Reference number: AB0502; A; Accession: AF0922
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  thioredoxin - Emericella nidulans
C;Species: Emericella nidulans, Aspergillus nidulans
C;Date: 22-Nov-1993 #sequence_revision 01-Sep-1995 #text_change 09-Jul-2004
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Best Local S
Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;2-109/Product: thioredoxin #status experimental;11-94/Domain: thioredoxin homology <THR>
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R,Nierman, W.C.; Feldblyum, T.V.; Paulsen, I.T.; Nelson, K.E.B.; Laub, M.T.; DeBoy, R.T.; Dodson, R.J.; Durkin, A.S.; Gwn, J.; Ermolaeva, M.; White, O.; Salzberg, S.L.; Shapiro, L. Proc. Natl. Acad. Sci. U.S.A. 98, 4136-4141, 2001
A;Title: Complete Genome Sequence of Callobacter crescentus. A;Reference number: A87249; MUID:21173698; PMID:11259647
A;Accession: A87688
                                                                                                                                                    A;Authors: Grandi, G.; Sun, L.; Smith, H.O.; Fraser, C.M.; Moxon, E.R.; A;Title: Complete genome sequence of Neisseria meningitidis serogroup B A;Reference number: A81000; MUID:20175755; PMID:10710307 A;Accession: C81090 A;Accession: C81090 A;Status: preliminary A;Molecule type: DNA A;Residues: 1-110 <TET>
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C; Superfamily:
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A;Molecule type: DNA
A;Residues: 1-110 <STO>
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C;Date: 20-Apr-2001 #sequence_revision 20-Apr-2001 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Gene: STY3639
C; Superfamily: thioredoxin; thioredoxin homology
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                                                          A;Gene: NMB1366
C;Superfamily:
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A;Experimental source: serogroup B, strain MC58
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strain MCS
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aft, D.H.; Kolon
C.; Fraser, C.M.
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A;Cross-references: UNIPARC:UPI000016DD37; EMBL:X78822; NID:g840740; PIDN:CAA55399.1; R;Decottignies, P.; Schmitter, J.M.; Dutka, S.; Jacquot, J.P.; Miginiac-Maslow, M. Bur. J. Biochem. 198, 505-512, 1991

A;Title: Characterization and primary structure of a second thioredoxin from the greer A;Reference number: S16090; MUID:91249849; PMID:2040309
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R;Stein, M.; Hodges, M.; Jeanette, E.; Lancelin, J.M.; Jacquot, submitted to the EMBL Data Library, April 1994
A;Description: Chlamydomonas reinhardtii thioredoxins I : cDNA;
A;Reference number: S54844
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R;Parkhill, J.; Achtman, M.; James, K.D.; Bentley, S.D.; Churcher, C.; Klee, S.R.; Morel ; Holroyd, S.; Jagels, K.; Leather, S.; Moule, S.; Mungall, K.; Quail, M.A.; Rajandream, Nature 404, 502-506, 2000
                                  A;Cross-references: UNIPARC:UPI0000171E27
R;Mittard, V.; Blackledge, M.J.; Stein, M.; Jacquot, J.P.;
submitted to the Brookhaven Protein Data Bank, May 1996
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A;Reference number: A81775; MUID:20222556; PMID:10761919
A;Accession: E81850
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                                                                                                                                                                 A; Molecule type: protein A; Residues: 2-112 < MIG>
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A; Residues: 1-113 <STF>
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Reference number: S57774; MUID:95359406; PMID:7632918
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PDB:1TOF
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Pred. No. 1.1;
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                                                                          Marion,
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                                                                                                                                                                                                                                                                                                                          thioredoxin from the green
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                                                                          D.; Lancelin,
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                                                                                                                                                                                       R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Hc Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, 537-544, 1998
A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complet A;Reference number: A70500; MUID:98295987; PMID:9634230
A;Accession: B70851
                          A; Molecule type: DNA
A; Residues: 1-116 <COL>
A; Cross-references: UNI
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                                                                                                                                                    A; Status: preliminary; nucleic acid sequence not shown; translation not
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probable trxC protein - Mycobacterium tuberculosis (strain H37RV)
C;Species: Mycobacterium tuberculosis
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nces: UNIPROT:P52229; source: strain H37Rv

UNIPARC: UPI0000165202;

GB:AL021426;

GB:AL123456; NID

complete

genome

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shown

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RESULT 43
T29044
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A;Title: (1)H, (13)C, (15)N-NMR resonance assignments of oxidized thioredoxin R spectroscopy and computer-assisted backbone assignment.
A;Reference number: A58618; MUID:95262711; PMID:7744070
                                                                                                                                                                      A; Introns:
C; Superfam:
                                                                                                                                                                                                              A;Gene: CESP:B0228.5
                                                                                                                                                                                                                                                      A;Cross-references: UNIPROT:Q09433; UNIPARC:UPI0000136D67; EMBL:U23168; PIDN:AAC38808.1; A;Experimental source: strain Bristol N2
                                                                                                                                                                                                                                                                                                                      A; Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                  A; Description: The sequenc A; Reference number: Z18324
                                                                                                                                                                                                                                                                                                                                                                                                      submitted to the EMBL Data Library, March A;Description: The sequence of C. elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Leimbach,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hypothetical protein 80228.5 - Caenorhabditis elegans C;Species: Caenorhabditis elegans
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                                                                                                                                                                                                                                                                                                                                                              A; Accession: T29044
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                                                                                                                                                                    Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;2-113/Product: thioredoxin h #status experimental;15-98/Domain: thioredoxin homology <THR>
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                                                                                    Matches
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Best Local Similarity
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                                                                                  Similarity 6; Conserv
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                                          WCGPCK 6
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  43
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                                                                                                      Score 47; DB Pred. No. 1.1;
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Pred. No. 1.1;
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cosmid B0228
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thioredoxin (trxA) homolog - Lyme disease spirochete C;Species: Borrelia burgdorferi (Lyme disease spirochete) C;Date: 13-Feb-1998 #sequence_revision 13-Feb-1998 #text_change 05-Oct-2004 C;Accession: E70107 R;Fraser, C.M.; Casjens, S.; Huang, W.M.; Sutton, G.G.; Clayton, R.; Lathigra, R.; Son, D.; Peterson, J.; Kerlavage, A.R.; Quackenbush, J.; Salzberg, S.; Hanson, M.; Bowman, C.; Garland, S.; Fujii, C.; Cotton, M.D.; Horst, K.; Roberts, K.; Hatch, Pature 390, 580-586, 1997 Nature 390, 580-586, 1997 A;Authors: Smith, H.O.; Venter, J.C. A;Title: Genomic sequence of a Lyme disease spirochaete, Borrelia burgdorferi. A;Reference number: A70100; MUID:98065943; PMID:9403685
                                                                                                                                                                                                                                                                                                                                          C;Accession: S34812
R;Brugidou, C.; Marty, I.; Chartier, Y.; Meyer, Y.
Mol. Gen. Genet. 238, 285-293, 1993
A;Title: The Nicotiana tabacum genome encodes two cytoplasmic A;Reference number: S34812; MUID:93241165; PMID:8479434
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A;Gene: trxC
C;Superfamily: thioredoxin; thioredoxin homology
F;15-98/Domain: thioredoxin homology <TXN>
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C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change
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E70107
RESULT
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A; Residues: 1-118 < BRU>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 46
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A; Residues: 1-117 < KLE>
                                                                                                                                                                                                                                                                                                                            A; Accession: S3481
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                                                                                                                                                                                                                                                                   Cross-references: UNIPROT: Q07090; UNIPARC: UPI0000136D63;
                                                                                                                                                                                                                  Superfamily:
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47
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                                                   WCGPCK 43
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Pred. No.
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A;Accession: T39387
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A; Residues: 1-127 <FEL>
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A; Introns: 33/2
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                                                                                   gasawara, N.; Yasunaga,
                                                                                                    R;Hayashi,
                                                                                                                                                                                                                RESULT 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Accession: S19498
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                                                                                                                                                                                                 B91218
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Best Local S
Matches 6
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thioredoxin ii - fission yeast (Schizosaccharomyces C;Species: Schizosaccharomyces pombe C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #C;Accession: T39387 R;Lyne, M.; Rajandream, M.A.; Barrell, B.G.; Xiang, submitted to the EMBL Data Library, January 1999
                                                                                                                                                                                                           A;Cross-references: UNIPROT:094504; UNIPARC:UPI000006BF82; EMBL:AL035085; PIDN:CAA22681. A;Experimental source: strain 972h-; cosmid c12D12
                                                                                         C; Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                  A;Status: preliminary; translated from GB/EMBL/DDBJ
Conservative
                        100.0%;
  <u>,</u>
                     Score 47; DB Pred. No. 1.2;
  Mismatches
                                             <u>ب</u>
  0
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                                                                                                                                                                                                                                                                                                                                                                                              Z.; Aves,
                                             Length 121
  <u>.</u>
  Gaps
  0
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thioredoxin homolog YCR083w - yeast (Saccharomyces cerevisiae) C;Species: Saccharomyces cerevisiae #sequence_revision Н 10-Sep-1999 #text_change 05-Oct-2004

Database, March 1992

A; Gene: SGD: TRX3; MIPS: YCR083w A;Cross-references: UNIPROT:P25372; UNIPARC: UPI0000137687; EMBL: X59720; NID: g1907116; P1

A;Map position: 3R C;Superfamily: thioredoxin; thioredoxin homology C;Keywords: redox-active disulfide E;34-115/Domain: thioredoxin homology <THR> A;Cross-references: SGD:S0000679 A;Map position: 3R ;55-58/Disulfide bonds: redox-active #status predicted 100.0%; Score 47; DB : Pred. No. 1.2; DB 1; Length 127;

0

Mismatches

0

Indels

<u>;</u>

0

C;Species: Escherichia coli C;Date: 18-Jul-2001 #sequence_revision 18-Jul-2001 #text_change C;Accession: B91218 thioredoxin 1 [imported] - Escherichia coli (strain O157:H7, substrain RIMD 0509952)

T.; Makino, K.; Ohnishi, M.; Kurokawa, K.; Ishii, K.; N.; Yasunaga, T.; Kuhara, S.; Shiba, T.; Hattori, M.; S Shinagawa, Yokoyama, K.; Shinagawa, H.

O

geno

DNA Res. 8, 11-22, 2001
A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli 0157:H7 and A;Reference number: A99629; MUID:21156231; PMID:11258796
A;Accession: B91218
A;Status: preliminary

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R;Theologis, A; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alon Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, ansen, N.F.; Hughes, B.; Huizar, L.
Nature 408, 816-820, 2000

A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, B.; Kim, C.A.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Marzia Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Tallo ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A;Accession: B96621
                                                                                                                                                                                                                                                                                                                                                                                                                                              probable thioredoxin F23H11.5 [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 05-Oct-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, D.J.; iller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta, E.; Potamousis, K.; A, Nature 409, 529-533, 2001
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                                                                                                        A; Molecule type: DNA
A; Residues: 1-129 <STO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Gene: trxA
C; Superfamily: thioredoxin; thioredoxin homology
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A;Residues: 1-127 <STO>
A;Cross-references: UNIPARC:UPI0000165538; GB:AE005174;
A;Experimental source: strain 0157:H7, substrain EDL933
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C; Superfamily:
                                            A;Gene: F23H11.5
                                                                C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                              C; Accession: B96621
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                                                                                    A;Cross-references: UNIPROT:Q9XIF4; UNIPARC:UPI000000BF0A; GB:AE005173; NID:g5080807;
                                                                                                                                                   A; Status: preliminary
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Best Local
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Pred. No. 1.
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Pred. No. 1.2;
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ewar, K.;
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                                                                                      RESULT 54
F97361
thioredoxin c-1 [imported] - Agrobacterium tumefaciens (strain C58, Cereon) C;Species: Agrobacterium tumefaciens C;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 05-Oct-2004 C;Accession: F97361
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RESULT 53
D71707
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T08084
                                                                                                                                          A;Gene: trxA; RFUUZ
C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                          A;Cross-references: UNIPARC:UPI0000164C06; GB:AJ235270; A;Experimental source: strain Madrid E C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:Q39591; UNIPARC:UPI0000129A19; EMBL:U43609; NID:g1163185; A;Experimental source: strain 1132D C;Comment: Seems to be associated circles.
                                                                                                                                                                                                                                                                                                                                  C;Accession: D71707

C;Accession: D71707

R;Andersson, S.G.E.; Zomorodipour, A.; Andersson, J.O.; Sicheritz-Ponten, T.; Alsmark, Nature 396, 133-140, 1998

A;Title: The genome sequence of Rickettsia prowazekii and the origin of mitochondria. A;Reference number: A71630; MUID:99039499; PMID:9823893

A;Accession: D71707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Patel-King, R.S.; Benashaki, S.E.; Harrison, A.; King, S.M.
J. Biol. Chem. 271, 6283-6291, 1996
A;Title: Two functional thioredoxins containg redox-senesitive vicinal dithiols from the A;Reference number: 216342; MUID:96198089; PMID:8626422
A;Accession: T08084
                                                                                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-130 < AND >
                                                                                                                                                                                                                                                                                                              A;Status: preliminary; nucleic acid sequence not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Species: Rickettsia prowazekii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           thioredoxin (trxA) RP002 - Rickettsia prowazekii
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N;Alternate names: thioredoxin homolog
C;Species: Chlamydomonas reinhardtii
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Best Local S
Matches 6
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                                                                               Similarity 6; Conser
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                                       WCGPCK 6
                                                                             100.0%; Score 47; DB 2; Length 130; llarity 100.0%; Pred. No. 1.2; Conservative 0; Mismatches 0; Indels
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Pred. No. 1.2;
Nismatches
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C; Date: V4-Fra. C.; Conster V4-Fra. C.; Caccession: B96721

R; Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaui, S.; Fisheologis, A.; Ecker, J.R.; Conway, A.B.; Conway, A.R.; Creasy, T. Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T. Chin, C.W.; Chung, M.; Huizar, L. Nature 408, 816-820, 2000

Nature 408, 816-820, 2000

Nature 408, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-820, 15-
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R;Lim, C.J.; Sa, J.H.; Fuchs, J.A.
Biochim. Biophys. Acta 1307, 13-16, 1996
A;Title: Identification of a third thioredoxin gene from Corynebacterium nephridii. A;Reference number: $70356; MUID:96254056; PMID:8652661
A;Accession: $70356
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A;Map position: circular chromosome
C;Superfamily: thioredoxin; thioredoxin homology
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A.; Liu, F.; Wollam, C.; Allinger, M.;
Science 294, 2323-2328, 2001
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A;Residues: 1-133 <KUR>
A;Cross-references: UNIPROT:Q8UJA6; UNIPARC:UPI00000D1691; GB:AE007869;
   A;Cross-references:
                                A; Molecule type:
A; Residues: 1-148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probable thioredoxin T17F3.9 [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 05-Oct-2004
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Residues: 1-145 <LIM>
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                                    1-148 <STO:
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UNIPROT:Q9CAS1; UNIPARC:UPI000000C0F3; GB:AE005173; NID:g6358805;
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100.0%; Pred. No. 1.3;
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB
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.S.; Maiti, R.;
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RESULT 58
T09495
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A;Gene: T17F3.9
A;Map position: 1
C;Superfamily: thioredoxin
                                                                                                                                                                                                                                                           A; Molecule type: mRNA
A: Residues: 1-177 < DUE>
                                                                                                                                                                                                                                                                                                                                  R;Duek, P.D.; Mora Garcia, S.; Wolosiuk, R.A. submitted to the EMBL Data Library, October 1 A;Reference number: Z16696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPROT:Q39592; UNIPARC:UPI0000129A1A;
A;Experimental source: strain 1132D
C;Comment: Seems to interact directly with the a dynein hea
C;Superfamily: thioredoxin; thioredoxin homology
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A;Title: Two functional thioredoxins containg redox-senesitive
A;Reference number: Z16342; MUID:96198089; PMID:8626422
A;Accession: T08086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    dynein light chain, 16K, outer arm - Chlamydomonas reinhardtii
N;Alternate names: thioredoxin homolog
C;Species: Chlamydomonas reinhardtii
                                                                                                                                         C;Superfamily: thioredoxin; thioredoxin home C;Keywords: chloroplast; redox-active disulf;79-162/Domain: thioredoxin homology <TXN>
                                                                                                                                                                                                                                                                                                               A; Reference number: A; Accession: T09495
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A; Residues: 1-156 < PAT>
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J. Biol. Chem. 271, 6283-6291, 1996
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C;Accession: T08086
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C;Date: 20-Sep-1999 #sequence_revision 20-Sep-1999
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                                                                                                                                                                                                       A; Genome: chloroplast
                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:Q9XGS0; UNIPARC:UPI000009E914;
                                                                                                                                                                                                                                                                                              A;Status: translated from GB/EMBL/DDB
                                                                                                                                                                                                                                                                                                                                                                                                                                               thioredoxin m - rape chloroplast
                                                                                                                                                                Superfamily: thioredoxin; thioredoxin homology; Superfamily: thioredoxin; thioredoxin homology; Keywords: chloroplast; redox-active disulfide
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0; Mismatches
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Pred. No. 1.5
); Mismatches
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C;Comment: This enzyme has amino-terminal heterogeneity. Thioredoxin m activates malate C;Comment: Arg-145 and Gly-164 are both essential for thioredoxin to interact with thior C;Superfamily: thioredoxin; thioredoxin homology C;Keywords: chloroplast; redox-active disulfide

F;1-67/Domain: transit peptide (chloroplast) #status predicted <TNP>
F;68-171/Product: thioredoxin mb #status experimental <PRB>
F;69-171/Product: thioredoxin md #status experimental <PRD>
F;09-171/Product: thioredoxin md #status experimental <PRD>
F;82-165/Domain: thioredoxin homology <THR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross references: UNIPROT:D07591; UNIPARC:UDI0000136B67; GB:X51462; GB:S84848; NID:g21 R;Maeda, K.; Tsugita, A.; Dalzoppo, D.; Vilbois, F.; Schurmann, P. Eur. J. Biochem. 154, 197-203, 1986
A;Title: Further characterization and amino acid sequence of m-type thioredoxins from sp A;Reference number: A91158; MUID:86108311; PMID:3510868
A;Accession: JT0023
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A; Residues: 68-82, 'G', 84-89, 'Q', 91, 'SE', 94, 'S', 96-127, 'T', 129-164, 'D', 166, 'S', 168, 'YQ'
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A; Accession: T00893
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Pred. No. 1.5
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hypothetical protein F14I3.8 [imported] - Arabidopsis thaliana (Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change C;Accession: E96539 R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kangarana (Mariana) Response for the control of th
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A;Title: Isolation, characterization and nucleotide sequence of a full-length pea cD A;Reference number: $20929; MUID:92256804; PMID:1581563
A;Recession: $20929
A;Molecule type: mRNA
A;Residues: 1-182 <LEP-
A;Cross-references: UNIPROT:P29450; UNIPARC:UPI0000136DE9; EMBL:X63537; NID:g20906; C;Keywords: chloredoxin; thioredoxin homology C;Keywords: chloroplast; redox-active disulfide P;1-74/Domain: transit peptide (chloroplast) #status predicted <TNP-
F;83-167/Domain: thioredoxin homology <THR-
F;106-109/Disulfide bonds: redox-active #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Mora-Garcia, S.B.F.; Rodriguez-Suarez, R.J.; Wolosiuk, R.A. submitted to the EMBL Data Library, August 1997 A;Reference number: Z16164 A;Accession: T07837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Lepiniec, L.; Hodges, M.; Gadal, P.; Cretin, Plant Mol. Biol. 18, 1023-1025, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       thioredoxin f precursor - garden pea
C;Species: Pisum sativum (garden pea)
C;Date: 12-Feb-1993 #sequence revision 12-Feb-1993 #text_change 09-Jul-2004
C;Accession: S20929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Description: activates fructose 1,6-bisphosphatase C;Superfamily: thioredoxin; thioredoxin homology C;Keywords: chloroplast; redox-active disulfide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Gene: TRXF
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C;Date: 14-May-1999 #sequence_revision 14-May-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  thioredoxin f precursor -
E96539
                  RESULT 63
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                                                                                                                              WCGPCK 6
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Pred. No. 1.6;
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A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White,

#text_change 05-Oct-2004

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Alonso,

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Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H. ansen, N.F.; Hughes, B.; Huzzar, L.
Nature 408, 816-820, 2010

A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R. Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A;Reference number: A86141; MUID:21016719; PMID:11130712
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A;Accession: F85044
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-186 <ST
                                                                                        R;annnymous, The European Union Arabidopsis Genome Sequencir
Nature 402, 769-777, 1999
A;Title: Sequence and analysis of chromosome 4 of the plant
A;Reference number: A85001; MUID:20083488; PMID:10617198
                                                                                                                                                                                probable M-type thioredoxin [imported] - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_chancC;Accession: F85044
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A;Residues: 1-185 <KUR>
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Best Local S
Matches 6
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Best Local S
Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Species: Streptococcus pneumoniae;
;Date: 03-Aug-2001 #sequence_revision 03-Aug-2001 #text_change 09-Jul-2004
;Accession: D95115
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type: DNA : 1-186 <STO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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S.L.; Lewis, M.R.;
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                                                                                                                  Arabidopsis thaliana
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  A;Description: activates fructose 1,6-bisphosphatase during photosynthesis
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C;Accession: S04661; A20273
R;Kamo, M.; Tsugita, A.; Wiessner, C.; Wedel, N.; Bartling, D.; Herrmann, R.G.; Aguilar, R;Kamo, M.; Tsugita, A.; Wiessner, C.; Wedel, N.; Bartling, D.; Herrmann, R.G.; Aguilar, Bur. J. Biochem. 182, 315-322, 1989
A;Title: Primary structure of spinach-chloroplast thioredoxin f. Protein sequencing and A; Accession: S04661
A;Molecule type: mRNA
A;Residues: 1-190 <KAM>
A;Residues: 1-190 <KAM>
A;Residues: 1-190 <KAM>
A;Cross-references: UNIPACT:P09856; UNIPARC:UPI0000136DEA; EMBL:X14959; NID:g21343; PIDN A;Cross-references: UNIPACT:P09856; UNIPARC:UPI0000136DEA; EMBL:X14959; NID:g21343; PIDN A;Residues: 10f. 110; Schuermann, P.
Biochem. Biophys. Res. Commun. 115, 1-7, 1983
Biochem. Biophys. Res. Commun. 115, 1-7, 1983
A;Title: Spinach chloroplast thioredoxins in evolutionary drift.
A;Reference number: A20273; MUID:83308759; PMID:6351859
A;Recidues: 106-119; NGDKEATCHLGVQQA',120 <TSU>
A;Residues: 106-119; NGDKEATCHLGVQQA',120 <TSU>
A;Residues: 106-119; NGDKEATCHLGVQQA',120 <TSU>
A;Residues: 106-119; NGDKEATCHLGVQQA',120 <TSU>
A;Comment: Thioredoxin f activates fructose 1,6-bisphosphatase during photosynthesis.
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: chloroplast; redox-active disulfide
F;1-17/Domain: transit peptide (chloroplast) #status predicted <TNP>
F;18-190/Product: thioredoxin f #status predicted <MAT>
F;191-175/Domain: thioredoxin f #status predicted <MAT>
F;191-175/Domain: thioredoxin f #status predicted <MAT>
Fill-117/Nichain: Transit peptide (chloroplast) #status predicted <MAT>
Fill-117/Nichain: Transit peptide (chloroplast) #status predicted <MAT>
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A;Gene: AT4903520
A;Map position: 4
C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                         R;Michalowski, C.B.; Bohnert, H.J.
submitted to the EMBL Data Library, June 1998
A;Description: A cDNA for thioredoxin F precursor from
                                                                                                                                                                                                                                                              thioredoxin f precursor - common ice plant C;Species: Mesembryanthemum crystallinum (common ic C;Date: 23-Jul-1999 #sequence_revision 23-Jul-1999 C;Accession: T12261
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C;Species: Spinacia oleracea (spinach)
C;Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change 09-Jul-2004
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                              A; Molecule type: mRNA
A; Residues: 1-191 < MIC>
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                                                                                    A; Status: preliminary; translated from GB/EMBL/DDB.
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A; Cross-references:
                                                                                                                   A; Accession: T12261
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   UNIPROT: 081332; UNIPARC: UPI0000136DE8; EMBL: AF069314; NID: g3202021;
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C;Superfamily: thioredoxin; thioredoxin homology C;Keywords: redox-active disulfide F;92-176/Domain: thioredoxin homology <TXN>
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                                                                                                                                                                                                                                                                                                                                                                             C;Species: Schizosaccharomyces pombe
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004
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J. Bacteriol. 183, 5709-5717, 2001
A;Authors: Yang, Y.; Young-Bellido, M.; Zhao, G.; Zook, C.; Baltz, R.H.; Jaskunas, A;Title: Genome of the Bacterium Streptococcus pneumoniae Strain R6.
A;Reference number: A97872; MUID:21429245; PMID:11544234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Hoskins, J.A.; Alborn Jr., W.; Arnold, J.; Blaszczak, L.; Burgett, S.; DeHoff, B. e, R.; LeBlanc, D.J.; Lee, L.N.; Lefkowitz, E.J.; Lu, J.; Matsushima, P.; McAhren, Y. P.; Sun, P.M.; Winkler, M.E.
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T40552
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A; Residues: 1-191 < KUR>
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                                                                                                                                                                                                            A; Experimental source:
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                                                                                                                                                                                                                                                A;Residues:
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                                                                                                                                                        Gene: SPAC577.08c
                                                                                                                                                                                                                                                             Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                        Accession: T40552
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Best Local Similarity
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Search completed: February 23, Job time : 32.5 secs

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RESULT 70

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thioredoxin trxA [imported] - Agrobacterium tumefaciens (strain C58, Dupont) (;Species: Agrobacterium tumefaciens C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 09-Jul-2004 C;Accession: AH3011 R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, Y.; Woo, lerage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, M.; McClel.; Karp, P.; Romero, P.; Zhang, S. Science 294, 2317-2323, 2001 A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.; Gordon-Kamm, ster, E.W. A;Fitle: The Genome of the Natural Genetic Engineer Agrobacterium tumefaciens C58. A;Reference number: AB2577; MUID:21608550; PMID:11743193
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                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-322 < KUR>
                                                                                                                                                                                              A; Map position: linear chromosome
                                                                                                                                                                                                                           A;Gene: trxA
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WCGPCK
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Q75CQ0 | ASHGO
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RESULT 1

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OR Tetradontoides
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A Vaillon O., Aury J.M., Brunet F., Petit J.L., Stange-Thomann N.,

Maucell E., Bouneau L., Fischer C., Ozouf-Costaz C., Bernot A.,

Nicaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,

A Nicaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,

A Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,

A Anthouard V., Jubin C., Cattolico L., Poulain J., De Berardinis V.,

Cruaud C., Duprat S., Brottier P., Coutanceau J.P., Gouzy J.,

A Cruaud C., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,

Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,

A Farra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,

Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,

A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,

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Eukaryota; Metazoa; Chordata; Cranitata; Vertebrata; Euteleostomi;

Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei;

Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;

Tetradontoidea; Tetraodontidae; Tetraodon.
                                                                                                                                                                                               Genoscope; Whitehead Institute Centre for Genome Research; Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases.
-i- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry whi
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13-SEP-2005 (TrEMBLrel. 31, Last sequence up
13-SEP-2005 (TrEMBLrel. 31, Last annotation
Chromosome undetermined SCAF8161, whole geno
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ORFNames=GSTENG00005813001;
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Q950IA5 EDOM
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Q950IA5 ENOM
Q5CUA5 SCHUA
Q5CUA5 SCHUA
Q5CUA5 MAIZE
Q75MO8 ORYSA
Q5CUA7 MAIZE
Q75MO8 TEMENI
Q4WUA ASPFU
Q50UA7 ENTHI
ERP38 NEUCR
Q91815 FUGRU
Q50UA7 MAGGR
Q91212 ORYSA
Q94212 ORYSA
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                       Score 47;
Pred. No.
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RESULT 3
QST936_HUMAN p
TD QST936_T
PEB-2005 (T
DT 01-FEB-2005 (T
DE Thioredoxin.
GN Name=TXN, ORFN
OS Homo sapiens (
OC Eukaryota; Mete
OC Mammalia; Euth
OC Momo. TaxID=960
RM (1]
CX NCBI_TAXID=960
RN (1]
CR WILLEOTIDE SEC
RA SUbmitted (MAY
DR SUBmitted (MAY
DR GO; GO:0006118
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Best Local S
Matches 6
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"An alternative splice variant of human the control of human the control
Williams S.;
Submitted (MAY-2005) to the E
EMBL; AL158158; CAI14067.1; -
SWR; Q57936; 1-85.
GO; GO:0005489; F:electron tr
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01-MAR-2004 (TrEMBLrel. 07, L
Thioredoxin delta 3 (Fragment)
Name=TXN delta 2
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PRINTS; PR00421; THIOREDOXIN,
PROSITE; PS00194; THIOREDOXIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       060744 HUMAN
060744;
                                                                                                                                                                                                                 NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Redox-active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE
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HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WCGPCK 6
                                                                                                                                                                                                                                                                               Eutheria;
                                                                                                                                                                                                                                                                                                                                                            ORFNames=RP11-427L11.1-002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 1
84 AA; 9321 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE.
                                                                                                                                                                    SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                     (Human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ω
                                                                                                                                                                                                                                                                    Chordata; Craniata; Vertebra
comphontoglires; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Homir
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     transport; IEA
                                 transporter activity; IEA
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                                                                               EMBL/GenBank/DDBJ
-; Genomic_DNA.
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    818FB84BA86E5721 CRC64;
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n thioredoxin.";
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                                                                                                                                                                                                                                                                               Vertebrata; Euteleostomi;
rimates; Catarrhini; Hominidae;
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                                                                                                                 databases
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RESULT OF COLOR OF CO
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ID Q9
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Q9HMD0_HALSA
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Best Local S
Matches 6
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Best Local
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EMBL; AY596297; AAV46961.1; -; Genomic DNA.
GO; GO:0005489; F:electron transporter activi
GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR006662; Thiored. A
InterPro; IPR005746; Thioredoxin.
InterPro; IPR005746; Thioredox dom2.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                     01-MAR-2001 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
Thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR012335; ThioredoxIn-like.
Pfam; PF00085; ThioredoxIn; 1.
PRINTS; PR00421; THIOREDOXIN.
PROSITE; PS00194; THIOREDOXIN; 1.
Name=trxA2; OrderedLocusNames=VNG2600G; Halobacterium salinarium (Halobacterium halobium). Archaea; Euryarchaeota; Halobacteria; Halobacteriales;
                                                                                                           Q9HMD0;
Q9HMD0;
01-MAR-2001 (TrEMBLrel. 1
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PubMed-15520287; DOI=10.1101/gr.2700304;
Baliga H., Facciotti M.T., Pan M., Glusman G.,
Baliga H.S., Bhannon P., Chiu Y., Weng R.S., Gan R.R., Hung P.,
Date S.V., Marcotte E., Hood L., Ng W.V.;
"Genome sequence of Haloarcula marismortui: a halophilic archaeon from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Haloarcula marismortui (Halobacterium marismortui)
Archaea; Euryarchaeota; Halobacteria; Halobacteria
Halobacteriaceae; Haloarcula.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Name=trxA3; OrderedLocusNames=rrnAC2110;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-FEB-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Redox-active center.

Redox-active Sah; 9452 MW; 3CC6254BD6AlD66F CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=2238;
                                                                                                                                                                                                                                                                                                                                                                                                                                         TIGRFAMs; TIGR01068; thioredoxin;
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88 AA; 10011 MW;
                                                                                                                                                                                                                                                 WCGPCK 17
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                                                                         16,
26,
                                                                       Created)
Last sequence update)
Last annotation updat
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Pred. No. 2.1;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                           Score 47; DB Pred. No. 2.1;
                                                                                                                                                                                                                                                                                                                                                                                                     CFE9B8FAC1F69857 CRC64;
                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                     89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            activity; IEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88
                                                                                                                                                                                                                                                                                                                                                               DB 2; Length 88;
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SEQUENCE
 EMBL;
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                                                                             as
   U75512;
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NUCLECTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

RE STRAIN=NRC-1 / ATCC 700922 / JCM 11081;

RX MEDLINE=20504483; PubMed=11016590; DOI=10.1073/pnas.190337797;

RA Mg W.V., Kennedy S.P., Mahairas G.G., Berquist B., Pan M.,

RA Shukla H.D., Lasky S.R., Baliga N.S., Thorsson V., Sbrogna J.,

RA Shukla H.D., Lasky S.R., Baliga N.S., Thorsson V., Sbrogna J.,

RA Shukla H.D., Lasky S.R., Baliga N.S., Thorsson V., Sbrogna J.,

RA Swartzell S., Weir D., Hall J., Dahl T.A., Welti R., Goo Y.A.,

RA Swartzell S., Weir R., Cruz R., Danson M.J., Hough D.W.,

RA Maddocks D.G., Jablonski P.E., Krebs M.P., Augevine C.M., Dale H.,

RA Maddocks D.G., Jablonski P.E., Krebs M.P., Augevine C.M., Dale H.,

RA Isenbarger T.A., Peck R.F., Pohlschroder M., Spudich J.L., Jung K.-H.,

RA Isenbarger T.A., Liang P., Riley M., Hood L., DasSarma S.;

REMBL; AE005133; AAG20641.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                15-JUL-1998
15-JUL-1998
13-SEP-2005
Thioredoxin
                                                                                                                                                                                                                                                         Haverkamp T., Schwenn J.D.; Submitted (AUG-1996) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=1058;
                                                                                                                                                                                                                                                                                                                                                                              Thiocapsa roseopersicina.
Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                    Name=trxA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     THIO_THIRO P96132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR006663; Thioredox_dom2
Pfam; PF00085; Thioredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GO; GO:0005489; F:electron transporter activity; IEA
GO; GO:0006118; F:electron transport; IEA.
GO; GO:0006810; F:transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR005746; Thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PIR; E84409; E84409.
HSSP; P10599; 1ERV.
                                                                                                                     This Swiss-Prot
                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE [GENOMIC DNA].
                                                                                                                                                                                                                                                                                                                                                               Chromatiaceae; Thiocapsa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TIGRFAMs; TIGR01068; thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Halobacteriaceae;
                                                                                                                                                  similarity).
SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                                                                    FUNCTION: Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulfide and catalyzes dithiol-disulfide exchange reactions (By
                                                                Long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PR00421; THIOREDOXIN.
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89 AA;
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(Rel.
(Rel.
(TRX)
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                                                                                   Swiss Institute of Bioinformatics Bioinformatics Institute. There a
AAB36882.1;
2TRX.
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                                                                                                                 entry is copyright. It is produced through
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36, Last sequence 48, Last annotation
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Pred. No. 2.2;
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RESULT 8
Q4Z518 PLABE
ID Q4Z518_PLABE PRELIMINARY;
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Best Local S
Matches 6
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Q4XKV3;
13-SEP-2005
13-SEP-2005
13-SEP-2005
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13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation updat
Thioredoxin, putative (Fragment).

ORFNames=PC300021.00.0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             preliminary data.

EMBL; CAAJ01005024; CAH82459.1; -; Genomic_DNA.

InterPro; IPR006662; Thiored.

InterPro; IPR006663; Thioredox dom2.

InterPro; IPR012336; Thioredoxin-like.

InterPro; IPR012335; Thioredoxin fold.

Pfam; PP00085; Thioredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos F. Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S., "A comprehensive survey of the Plasmodium life cycle by genomic, transcriptomic, and proteomic analyses."; Science 307:82-86(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A., Berriman M., Florens L., Janssen C.S., Pain A., Christophides James K., Rutherford K., Harris B., Harris D., Churcher C., Quail M.A., Ormond D., Dogett J., Trueman H.E., Mendoza J., Bridell R. B., Berlander G., Dogett J., Trueman H.E., Mendoza J.,
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Hall N., Karras M., Raine J.D.,
Berriman M., Florens L., Jansse
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; ThioredoxIn.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
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Eukaryota, Alveolata, Apicomplexa, Haemosporida, Plasmodium
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PROSITE; PS00194; THIOREDOXIN;
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TIGRFAMs; TIGR01068; thioredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                           Redox-active center.
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13-SEP-2005 (TrEMBLrel.
13-SEP-2005 (TrEMBLrel.
13-SEP-2005 (TrEMBLrel.
Thioredoxin, putative (F.
ORFNames=PB000593,00.0;
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                                                                                                            gene expression.";
Science 307:1463-1465(2005).
EMBL; CR626927; CAH08710.1; -;
                                                                                                                                                                                       NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

PubMed=15746427; DOI=10.1126/science.1107008;

Cerdeno-Tarraga A.-M., Patrick S., Crossman L.C., Blakely G.,

Abratt V., Lennard N., Poxton I., Duerden B., Harris B., Quai

Barron A., Clark L., Corton C., Doggett J., Holden M.T.G., La

Line A., Lord A., Norbertczak H., Ormond D., Price C.,

Rabbinowitesh E., Woodward J., Barrell B.G., Parkhill J.;

"Extensive DNA inversions in the B. fragilis genome control v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
PROSITE; PS00194; THIOREDOXIN; UNKNOWN_1.
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EMBL; CAAIO1000448; CAH94443.1; -; Genc
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
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Eukaryota; Alveolata;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteroides fragilis (strain ATCC 25285
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NON_TER 1
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Hall N., Karras M.,
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                                    IPR006662; Thiored.
IPR006663; Thioredox dom2.
IPR005746; Thioredoxin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96 AA; 10769
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Last annotation updat
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           / NCTC 9343).
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RESULT 11
QBAB91 BACTN ID QBAB91;
AC QBAB91;
AC QBAB91;
DT 01-UUN-2003 (7)
DT 01-MAR-2004 (7)
DE Thioredoxin C-GN OrderedLocusNe
OS Bacteroides th
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DT 25-OCT-2

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PubMed=15466707; DOI=10.1073/pnas.0404172101;
Kuwahara T., Yamashita A., Hirakawa H., Nakayama H., Toh H.,
Kuhara S., Hattori M., Hayashi T., Ohnishi Y.;
"Genomic analysis of Bacteroides fragilis reveals extensive inversions regulating cell surface adaptation.";
Proc. Natl. Acad. Sci. U.S.A. 101:14919-14924(2004).
EMBL; AP006841; BAD49920.1; -; Genomic DNA.
GO; GO:000518; F:electron transporter activity; IEA.
GO; GO:000618; P:electron transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
                    OrderedLocusNames=BT0219;
                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                           TIGRFAMS; TIGR01068; thioredoxin; 1. PROSITE; PS00194; THIOREDOXIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bacteroidaceae; Bacteroides.
NCBI_TaxID=817;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bacteria; Bacteroidetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bacteroides fragilis.
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TIGRFAMS; TIGR01068; thioredo
PROSITE, P800194, THIOREDOXIN
Complete proteome.
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Pred. No. 2.3;
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RESULT 12
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GO; GO:0006118; P:electron transport; IEA.
GO; GO:0006810; P:transport; IEA.
InterPro; IPR006452; SH3.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredoxin.
InterPro; IPR006563; Thioredoxin.
InterPro; IPR006663; Thioredoxin.
InterPro; IPR006663; Thioredoxin.
InterPro; IPR006563; Thioredoxin; 1.
PRINTS; PR00452; SH3DOMAIN.
PRINTS; PR00452; SH3DOMAIN.
PRINTS; PR00452; THIOREDOXIN.
ITIGRPAMS; TIGR01068; thioredoxin; 1.
PROSITE; PR00154; THIOREDOXIN; 1.
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10-OCT-2003 (Rel. 4
10-MAY-2005 (Rel. 4
Thioredoxin (Trx).
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MEDLINE=21956528; PubMed=11959450;
Skamrov A.V., Feoktistova E.S., Gol'dman M.A.,
"Mycoplasma gallisepticum rpoA gene cluster.";
FEMS Microbiol. Lett. 208:281-285(2002).
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                                                                                                                          Microbiology 149:2307-2316(2003).
-!- FUNCTION: Participates in various redox reversible oxidation of its active center and catalyzes dithiol-disulfide exchange
                                                                                                                                                                                                                                                                       "The complete genome sequence gallisepticum strain R(low).";
                                                                                                                                                                                                                                                                                                            Papazisi L., Gorton T.S., Kutish G., Markha
Nguyen D.K., Swartzell S., Madan A., Mahair
"The complete genome sequence of the avian
                                                                                                                                                                                                                                                                                                                                             MEDLINE=22830409; PubMed=12949158; DOI=10.1099/mic.0.26427-0;
Papazisi L., Gorton T.S., Kutish G., Markham P.F., Browning G
Nguyen D.K., Swartzell S., Madan A., Mahairas G., Geary S.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=trxA; Synonyms=trx; OrderedLocusNames=MYCGA6330; OrFNames=MGA_0452; Mycoplasma gallisepticum.
Bacteria; Firmicutes; Mollicutes; Mycoplasmataceae; Mycop
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"A genomic view of the human-Bacteroides thetaiotaomicron symbiosis.";
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STRAIN=VPI-5482 / AT
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                             SIMILARITY: Belongs to the thioredoxin family. SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                       similarity)
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PubMed=12663928; DOI=10.1126/science.1080029;
PubMed=17663928; Deng S., Carmichael L.K.,
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Pred. No. 2.4;
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RESULT 13
Q9JN64 MYCGA
ID Q9JN64;
AC Q9JN64;
AC Q9JN64;
DT 01-CCT-2000 (T
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DT 01-MAR-2004 (T
DT 01-MAR-2009
RN NUCLEOTIDE SEQ
RC STRAIN=8596;
RX MEDLINE=963200
RA Skamrov A.V.
RT PEBS Lett. 291
RN NUCLEOTIDE SEQ
RC STRAIN=8596;
RX MEDLINE=963200
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                Skamrov A., Feoktistova E., (
Submitted (PEB-2003) to the I
EMBL; L35043; AAF36768.1; -;
HSSP; P80579; 1NW2.
                                                                                                                            MEDLINE=96320025; PubMed=8754006; Skamrov A.V., Bibilashvili R.Sh.; Skamrov alv., Bibilashvili R.Sh.; Paphysical map of Mycoplasma gallisepticum strain determination of its positions on certain genes."; Mol. Biol. (Mosk.) 30:585-594(1996).
                                                                                                                                                                                                                                                                                       Skamrov A.V., Bibilashvili R.S.; "Mycoplasma gallisepticum strain S6 genome contains three regions hybridizing with 16 S rRNA and two regions hybridizing with 23 S
                                                                                                                                                                                                                                                                                                                            STRAIN=A5969;
MEDLINE=92038016; PubMed=1718781; DOI=10.1016/0014-5793(91)81106-I;
                                                                                                                                                                                                                                                                                                                                                                                                                   Mycoplasma gallisepticum.
Bacteria; Firmicutes; Mollicutes;
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PRINTS; PR00421; THIOREDOXIN.
ProDom; PD003679; Thioredoxin_like;
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InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-1ike.
InterPro; IPR012335; Thioredoxin fold.
InterPro; IPR011594; Thioredoxin—like.
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100 AA;
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9; AAP56983.1; ALT_INIT; Genomic_DNA
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                                               Goldman M., Beabealashvilli R.; EMBL/GenBank/DDBJ databases.
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Pred. No. 2.4;
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A Dodson R.J., DeBoy R.T., Gwinn M.L., Nelson W.C., Haft D.H.,
A Hickey E.K., Peterson J.D., Durkin A.S., Kolonay J.F., Yang F.,
A Holt I.E., Umayam L.A., Mason T.M., Brenner M., Shea T.P.,
A Holt I.E., Umayam L.A., Mason T.M., Brenner M., Shea T.P.,
A Radune D., Vamathevan J.J., Khouri H.M., White O., Gruber T.M.,
A Ketchum K.A., Venter J.C., Tettelin H., Bryant D.A., Fraser C.M.;
The complete genome sequence of Chlorobium tepidum TLS, a
photosynthetic, anaerobic, green-sulfur bacterium.";
Proc. Natl. Acad. Sci. U.S.A. 99:9509-9514 (2002).

C -!- FUNCTION: Participates in various redox reactions through the
reversible oxidation of its active center dithiol to a disulfide
and catalyzes dithiol-disulfide exchange reactions.

-!- SIMILARITY: Belongs to the thioredoxin family.
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Best Local S
Matches 6
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13-SEP-2005 (Rel. 44, Last sequence uping-sep-2005 (Rel. 48, Last annotation Thioredoxin 1 (Trx-1).
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PROSITE; F
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InterPro;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chlorobium tepidum.
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PRINTS; PR00421; THIOREDOXIN
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n; PF000085; Thioredoxin; 1.
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DISULFID
SEQUENCE
                            Chlamydophila caviae. Bacteria; Chlamydiae;
                                                  Name=trxA; Synonyms=trx; OrderedLocusNames=CCA00080;
                                                             Thioredoxin
                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                   "The complete genome and proteome of Mycoplasma Genome Res. 14:1447-1461(2004).
EMBL; AE017337; AAR27629.1; -; Genomic_DNA.
HSSP; P10599; 1AUC.
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Butler J., Calvo S., Elkins T., FitzGerald M.G., Haf
Kodira C.D., Major J., Wang S., Wilkinson J., Nicol
Birren B., Berg H.C., Church G.M.;
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PubMed=15289470; DOI=10.1101/gr.2674004;
Jaffe J.D., Stange-Thomann N., Smith C., DeCaprio
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STRAIN=163K / ATCC
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                NCBI_TaxID=83557;
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16-OCT-2001 (Rel. 4
10-MAY-2005 (Rel. 4
Thioredoxin (TRX).
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Q9PJK3;
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DISULFID
SEQUENCE
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                                                                                             Chlamydia muridarum.
Bacteria; Chlamydiae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; L39892; AAB41348.1; -; Genomic_DNA.
EMBL; AE016994; AAP04832.1; -; Genomic_DN
HSSP; P80579; 1QUW.
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Microbiology 142:945-953(1996).
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TIGRFAM6; TIGR01068; thioredoxin; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
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STRAIN=MoPn / Nigg;
                       NUCLEOTIDE SEQUENCE
                                                                   NCBI_TaxID=83560;
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MEDLINE=97090413; PubMed=8936321;
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Nucleic Acids Res. 28:1397-1406(2000).

Nucleic Acids Res. 28:1397-1406(2000).
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                                                                                               Chlamydia trachomatis.";
Science 282:754-759(1998).
                                                                                                                                                                                       MEDLINE=99000809; PubMed=9784136; DOI=10.1126/science.282.5389.754; Stephens R.S., Kalman S., Lammel C.J., Fan J., Marathe R., Aravind Mitchell W.P., Olinger L., Tatusov R.L., Zhao Q., Koonin E.V.,
                                                                                                                                                                                                                                                                                                                                                                 Name=trxA; OrderedLocusNames=CT539;
Chlamydia trachomatis.
Bacteria; Chlamydiae; Chlamydiales; Chlamydiaceae; Chlamydia
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InterPro; IPR005746; ThioredoxIn.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
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                 FUNCTION: Participates in various redox reactions through reversible oxidation of its active center dithiol to a distant catalyzes dithiol-disulfide exchange reactions.
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28 31 Redox-active (By similarity).
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RESULT 19
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[2]
NUCLEOTIDE SEQUENCE OF 1-21.
STRAIN=ATCC 33530 / G-37;
STRAIN=94075230; PubMed=8253680;
MEDLINE=94075230, Hu P.-C., Bott K.F., Hut
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P47370; Q49453;
01-FEB-1996 (Rel
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                                                                                                                                                                                                 STRAIN=ATCC 33530 / G-37;

MEDLINE=96026346; PubMed=7569993;

Fraser C.M., Gocayne J.D., White O., Adams M.D., Clayton R.A., Fleischmann R.D., Bult C.J., Kerlavage A.R., Sutton G.G., Keil Fritchman J.L., Weidman J.F., Small K.V., Sandusky M., Fuhrmar Nguyen D.T., Utterback T.R., Saudek D.M., Phillips C.A., Merri Tomb J.-F., Dougherty B.A., Bott K.F., Hu P.-C., Lucier T.S., Peterson S.N., Smith H.O., Hutchison C.A. III, Venter J.C.; "The minimal gene complement of Mycoplasma genitalium."; Science 270:397-403(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR006663; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam, PF00085; Thioredoxin_fold.
                                               J. Bacteriol. 175:7918-7930(1993).
-i- FUNCTION: Participates in various reversable oxidation of its active and catalyzes dithiol-disulfide exc
                                                                                                                                                                                                                                                                                                                                                                            Name=trxA; Synonyms=trx; OrderedLocusNames=MG124; Mycoplasma genitalium. Bacteria; Firmicutes; Mollicutes; Mycoplasmatacea.
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10-MAY-2005 (Rel.
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PHCI-2DPAGE; 084544;
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         similarity).
SIMILARITY: Belongs to
SIMILARITY: Contains 1
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102 AA;
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31 Redox-active (By similarity).
11197 MW; BPAE022A5C980BE1 CRC64;
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Merrick J.M.,
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PROSITE; F
                           This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                               Nucleic Acids Res. 24:4420-4449(1996).
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MEDLINE-97105885; PubMed-8948633; DOI=10.1093/nar/24.22.4420;
Himmedreich R., Hilbert H., Plagens H., Pirkl E., Li B.-C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "The thioredoxin reductase system Microbiology 143:1933-1940(1997).
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Rottem S.;
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STRAIN=ATCC 29342 / M129;
MEDLINE=97346047; PubMed=9202470;
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Bacteria; Firmicutes; Mollicutes;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-11ke.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
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EMBL; U01796; AAD12321.1; ALT_INIT; Genomic_DNA.
PIR; G64213; G64213.
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                                                                                                                                                                                                                                                                                                                     'Complete sequence analysis of the genome of the
                                                                                                                                                                                          FUNCTION: Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulfide and catalyzes dithiol-disulfide exchange reactions (By
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B; TIGR01068; thioredoxin; 1.

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Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes; Schizosaccharomycetales; Schizosaccharomycetales; Schizosaccharomyces.
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
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Cho Y.-W., Shin Y.H., Kim
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GO; GO:0005737; C:cytoplasm; TAS.

GO; GO:0005737; C:cytoplasm; TAS.

GO; GO:000302; P:response to reactive oxygen specific properties of the control of the contro
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L_YEAST
TRX1_YEAST
P22217;
01-AUG-1991
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Thioredoxin
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TIGRFAMS; TIGR01068; thioredoxin; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
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SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; AF251279; AAF76881.1; -; Genomic_D; AF192765; AAF05765.1; -; mRNA.; AJ003819; CAA06033.1; -; mRNA.; Z99532; CAB16724.1; -; Genomic_DNA. T39085; T39085.
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 (Rel. 19, Created)
(Rel. 34, Last sequence update)
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I (TR-I) (Thioredoxin 2).
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Nucleophile (By similarity).
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Duesterhoeft A., Entian K.-D., Floeth M., Goffeau A., Hebling U., Heumann K., Heuss-Neitzel D., Hilbert H., Hilger F., Kleine K., Koetter P., Louis E.J., Messenguy F., Mewes H.-W., Miosga T., Koettler P., Louis E.J., Messenguy F., Mewes H.-W., Miosga T., Mosstl D., Mueller-Auer S., Nentwich U., Obermaier B., Piravandi Pohl T.M., Portetelle D., Purnelle B., Rechmann S., Rieger M., Rinke M., Rose M., Scharfe M., Scherens B., Scholler P., Schwager Schwarz S., Underwood A.P., Urrestarazu L.A., Vandenbol M., Verhasselt P., Vierendeels F., Voet M., Volckaert G., Voss H., Wambutt R., Wedler E., Wedler H., Zimmermann F.K., Zollner A., Vandenbol M.,
                                                                                                                                                                                                                                                                                  PROTEIN SEQUENCE OF 1-12, AND TSA1 DEPENDE
STRAINSATCC 200358 / YNN 295;
MEDLINE=95050519; PubMed=7961686;
Chae H.Z., Chung S.J., Rhee S.G.;
"Thioredoxin-dependent peroxide reductase
J. Biol. Chem. 269:27670-27678(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                         "Creation of the YFLEX clone resource: cloning cerevisiae ORFs in the Gateway recombinational Submitted (FEB-2004) to the EMBL/GenBank/DDBJ d
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Marsischky G., Rolfs A., Richardson A., Kane M., Baqui M., Taycher Hu Y., Vannberg F., Weger J., Kramer J., Moreira D., Kelley F., Zuo D., Raphael J., Hogle C., Jepson D., Williamson J., Camargo A., Gonzaga L., Vasconcelos A.T., Simpson A., Kolodner R., Harlow E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=97313267, PubMed=9169871,
Johnston M., Hillier L.W., Riles L.
Ansorge W., Benes V. Brucckner M.,
Duesterhoeft A., Entian K.-D., Floet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Muller E.G.D.;
"Thioredoxin genes in and TRX2.";
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MEDLINE=91225027; PubMed=2026619;
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MEDLINE=91107668; PubMed=1988444;
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Saccharomycetales; Saccharomycetaceae; Saccharomyces
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PubMed=1561834;
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                        Schwenn
                                                                                                                                                                                                           PROTEIN SEQUENCE OF 25-33.
MEDLINE=72104597; PubMed=4333602;
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J. Biol. Chem. 266:1692-1696(1991).
                                              PubMed=3060034;
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                                                                                                                   D.E., Baldesten A., Holmgren A., Reichard P.; covalently-bound flavin of hepatic monoamine oxidas tification and properties of cysteinyl riboflavin."; J. Biochem. 24:328-335(1971).
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nucleotide sequence
re 387:87-90(1997).
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  n J.D., Krone F. PAPS reductase:
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a., Hebling U.,
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Grant
"Role
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PubMed=9657146; DOI=10.1016/S0092-8674(00)81457-9;

Xu Z., Sato K., Wickner W.;

"LMA1 binds to vacuoles at Sec18p (NSF), transfers

to a t-SNARE (Vam3p) complex, and is released during the second of t
Ghaemmaghami S., Huh W.-K., Dephoure N., O'Shea E.K., We "Global analysis of protein Nature 425:737-741(2003).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNCTION, AND REGULATION OF HYR1/GPX3.
PUNMed=12437921; DOI=10.1016/S0092-8674(02)01048-6;
Delaunay A., Pfileger D., Barrault M.-B., Vinh J.,
"A thiol peroxidase is an H2O2 receptor and redox-t:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=99143107; PubMed=9988687; DOI=10.1074/jbc.274.8.4537; Lee J., Spector D., Godon C., Labarre J., Toledano M.B.; "A new antioxidant with alkyl hydroperoxide defense propertie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Distinct physiological functions of thiol peroxidase isoenzymes Saccharomyces cerevisiae.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  to promote yeast vacuole inheritance.";
J. Cell Biol. 136:299-306(1997).
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Xu Z., Mayer A., Muller E.G.D., Wickner W.;
"A heterodimer of thioredoxin and I(B)2 cooperates with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Annu. Rev. Microbiol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     activation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Thioredoxins are required for protection in the yeast Saccharomyces cerevisiae."; Mol. Microbiol. 46:869-878(2002).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Roles of the glutathione- and thioredoxin-dependent in the Escherichia coli and Saccharomyces cerevisiae
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                                                                                                                                                       Biophys. Acta 1641:145-156(2003)
                                                                                                                                                                                                                                                                                                                                                           the glutathione/glutaredoxin and thioredoxin systems
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RESULT 23
Q5L731 CHLAB
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CH Name=trx
OS Chlamydo
OC Bacteria
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Matches 6
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Q5L731;
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Q1-FEB-2005 (TrEMBLrel. 29, C
01-FEB-2005 (TrEMBLrel. 29, I
01-FEB-2005 (TrEMBLrel. 29, I
Putative thioredoxin.
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MEDLINE=22923954; PubMed=14562095; DOI=10.1038/nature02026;
Huh M.-K., Falvo J.V., Gerke L.C., Carroll A.S., Howson R.W
Weissman J.S., O'Shea E.K.;
"Global analysis of protein localization in budding yeast."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hall D.E., Baldesten A., Holmgren A., Reichard "Yeast thioredoxin. Amino-acid sequence around disulfide of thioredoxin I and II.";
Eur. J. Blochem. 23:328-335(1971).
                                                                                                                                                                                                                                                                                                                                                                                                                                use
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                                                                                          Name=trxA; OrderedLocusNames=CAB082
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hydroperoxide reductase DOT5, thioredoxin peroxidases TSA1 and TSA2, alkyl hydroperoxide reductase AHP1, and peroxiredoxin HYR1. Thioredoxin is also involved in protection against reducing stress. As part of the LMA1 complex, it is involved in the facilitation of vesicle fusion such as homotypic vacuole and ER-derived COPII vesicle fusion with the Golgi. This activity does not require the redox mechanism.

SUBUNIT: Part of the heterodimeric LMA1 complex together with the proteinase inhibitor PBI2. LMA1 binds to the ATPase SEC18.
                                                                                                                                                                                                                                                                                                                                                                                                                                              s Swiss-Prot entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the EN European Bioinformatics Institute. There are no resti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TRX2, and one mitochondrial, TRX3.
SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protons. It is involved in many cellular processes, i deoxyribonucleotide synthesis, repair of oxidatively proteins, protein folding, sulfur metabolism, and red homeostasis. Thioredoxin-dependent enzymes include phosphoadenosine-phosphosulfate reductase METI5, alang budgephotide reductase horistases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FUNCTION: Participates as a hydrogen donor in redox reactions through the reversible oxidation of its active center dithiol disulfide, accompanied by the transfer of 2 electrons and 2
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Yeast has two cytoplasmic thiored
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Chlamydophila abortus Bacteria; Chlamydiae;

Chlamydiae;

Chlamydiales;

Chlamydiaceae;

Chlamydophila

STRAIN=S26/3; NUCLEOTIDE NCBI_TaxID=83555;

SEQUENCE

[LARGE

SCALE

GENOMIC DNA]

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RESULT
Q6MUGO
Q6MUGO
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Best Local S
Matches 6
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GO; GO:0005489; F:electron transporter activity; GO; GO:0006118; F:electron transport; IEA.
GO; GO:0006810; P:transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR00663; Thioredox dom2.
Pfam; PF00085; Thioredoxin; 1.
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Q6MUGO;
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THIOREDOXIN.
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Thomson N.R., Yeats C., Bell K., Holden M.T.G., Bentley S.D.,
Livingstone M., Cerdeno-Tarraga A.-M., Harris B., Doggett J.,
Ormond D., Mungall K., Clarke K., Feltwell T., Hance Z., Sanders
Quail M.A., Price C., Barrell B.G., Parkhill J., Longbottom D.;
"The Chlamydophila abortus genome sequence reveals an array of
variable proteins that contribute to interspecies variation.";
Genome Res. 15:629-640(2005).
EMBL; CR848038; CAH63539.1; -; Genomic_DNA.
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Butler J., Elkins
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Bacteria; Firmicutes; Mollicutes; Entomo
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   Mycoplasma
                                    Name=trxA;
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ATCC 33453;
, Stange-Thomann N., Har Elkins T., Kodira C.D.,
   mycoides
                                    OrderedLocusNames=MSC
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D., Major J.,
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RESULT 26
THIO_BACSU
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Best Local S
Matches 6
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P14949; 007960; Q45687;
01-APR-1990 (Rel. 14, Creat
01-OCT-1996 (Rel. 34, Last
10-MAY-2005 (Rel. 47, Last
MEDLINE=98044033; PubMed=9384377; DOI=10.1038/36786;
Kunst F., Ogasawara N., Moszer I., Albertini A.M., Alloni G.,
Azevedo V., Bertero M.G., Bessieres P., Bolotin A., Borchert
Borriss R., Boursier L., Brans A., Brann M., Brignell S.C., E
Brouillet S., Bruschi C.V., Caldwell B., Capuano V., Carter N
                                                                                                                                                                    Emmerson P.T., Harwood C.R.;
"The dnaB-pheA (256 degrees-240 degrees) region of the Bacil subtilis chromosome containing genes responsible for stress the utilization of plant cell walls and primary metabolism." Microbiology 142:3067-3078(1996).
                                                                                                                                                                                                                                                                                                                                                                                                          Chen N.-Y., Zhang J.-J., Paulus H.; "Chromosomal location of the Bacillus subtilis aspartokinase II gene and nucleotide sequence of the adjacent genes homologous to uvrC and trx of Escherichia coli.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=trxA; Synonyms=trx; OrderedLocusNames=BSU28500;
Bacillus subtilis.
Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GO; GO:0005489; F:electron transporter activity; IEA. GO; GO:0006118; P:electron transport; IEA. GO; GO:0006810; P:transport; IEA. InterPro; IPR006662; Thiored. InterPro; IPR005746; Thioredoxin. InterPro; IPR005746; Thioredox dom2. Pfam; PF00085; Thioredoxin, 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Complete proteome. SEQUENCE 102 AA;
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HSSP; P10599; IAUC.
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NCBI_TaxID=44101;
                                                                                                                                                                                                                                                                           MEDLINE=97124191; PubMed=8969504; Wipat A., Carter N., Brignell C.S., Guy J.B., Piper K.,
                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE
STRAIN=168;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=90132525; PubMed=2559145;
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                                                                                                                                 [LARGE SCALE
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Pred. No. 2.4
0; Mismatches
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    Carter N.M.,
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RA BELLEGE F., Devile K.M., Duestelloelt M., Billian K.-D., Eminical S.J., Bulliet S.J., Bulliet S.J., Bulliet S.J., Bulliet S.J., Balleron N., RA Fritz C., Fujita M., Fujita Y., Fuma S., Galizzi A., Galleron N., RA Fritz C., Fujita M., Fujita Y., Fuma S., Galizzi A., Galleron N., RA Guiseppi G., Gyy B.J., Hosono S., Hullo M.-F., Itaya M., Jones L.-M., Joris B., Karamata D., Kasahara Y., Klaerr-Blanchard M., RA Hilbert H., Holsappel S., Hosono S., Hullo M.-F., Itaya M., Jones L.-M., Joris B., Karamata D., Kasahara Y., Klaerr-Blanchard M., RA Klein C., Kobayashi Y., Koetter P., Koningstein G., Krogh S., RA Klein C., Koyayashi Y., Koetter P., Koningstein G., Krogh S., RA Klein C., Kordina N., Hellado R.P., Mizuno M., Moseti D., Nakai S., RA Medigue C., Medina N., Mellado R.P., Mizuno M., Moseti D., Nakai S., RA Medigue C., Medina N., Mellado R.P., Mizuno M., Moseti D., Nakai S., RA Mošack M., Noone D., O'Reilly M., Ogawa K., Ogiwara A., Oudega B., RA Medigue C., Medina N., Persecan E., Pujic P., Purnelle B., Rapoport G., RA Prescott A.M., Presecan E., Pujic P., Purnelle B., Rapoport G., RA Park S.-H., Parro V., Pohl T.M., Porteelele D., Porvollik S., Raport G., RA Rey M., Reynolds S., Rieger M., Rivolta C., Rocha E., Roche B., Raport G., Schi S., Schroeter R., RA Rose M., Sadaie Y., Sato T., Scanlan E., Schleich S., Schroeter R., RA Rose M., Sadaie Y., Sato T., Scanlan E., Schleich S., Schroeter R., RA Rose M., Sadaie Y., Sato T., Scanlan E., Schleich S., Schroeter P., RA Tognoni A., Toatch V., Uchiyama S., Vandenbol M., Vannier F., RA Weitzenegger T., Winters P., Wipat A., Yamamoto H., Yamane K., RA Weitzenegger T., Winters P., Wipat A., Yamamoto H., Yamane K., RA Yasunoto K., Yata K., Yoshida K., Yoshikawa H.-F., Zumstein E., Rank Yoshikawa H.-F., Zumstein E.,
                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; J03294; AAA8731;
EMBL; Z75208; CAA9957
EMBL; Z99118; CAB1481,
EMBL; X79976; CAA5630;
EMBL; X99275; CAA6766
PIR; B37192; B37192.
HSSP; P23400; 1DBY.
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InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
InterPro; IPR011594; Thioredoxin like.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
ProDom; PB003679; Thioredoxin like; 1.
TIGRFAM8; TIGR01068; thioredoxin; 1.
PROSITS; PS00194; THIOREDOXIN; 1.
PROSITS; PS00194; THIOREDOXIN; 1.
Complete proteome; Direct protein sequencing; Electron transport; Redox-active center; Transport.
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Graumann P., Schroeder K., Schmid R.,
"Cold shock stress-induced proteins i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   reversible oxidation of its active center dithiol to a and catalyzes dithiol-disulfide exchange reactions. SIMILARITY: Belongs to the thioredoxin family. SIMILARITY: Contains 1 thioredoxin domain.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Swiss-Prot entry is copyright. It is produced through sen the Swiss Institute of Bioinformatics and the EN Buropean Bioinformatics Institute. There are no restr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            complete
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(NOV-1994) to the
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; CAA99577.1;
; CAB14810.1;
; CAA56300.1;
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EMBL/GenBank/DDBJ databases.
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n Bacillus subtilis.";
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RESULT 27
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SEQUENCE
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Arroyo J., Backes U., Barreiros T., Bertani I., Bjourson A.J.,
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Submitted
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01-AUG-1991
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                                                                                                                                                                                                           phosphoserine phosphohydrolase,
Yeast 12:273-280(1996)
[5]
                                                                                                                                                                                                                                                   "Sequencing of a 17.6 kb segment on the right arm of yeast chromoson VII reveals 12 ORFs, including CCT, ADES and TR-I genes, homologues the yeast PMT and EFIG genes, of the human and bacterial electrontransferring flavoproteins (beta-chain) and of the Escherichia coli
                                                                                                                                                                                                                                                                                                                       MEDLINE=97060019; PubMed=8904340;
DOI=10.1002/(SICI)1097-0061(19960315)12:3<273::AID-YEA898>3.3.CO;2-T;
Guerreiro P., Barreiros T., Soares H., Cyrne L., Maia e Silva A.,
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Chem. 266:9194-9202(1991)
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225027; PubMed=2026619;
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Marsischky G., Rolfs A., Richardson A., Kane M., Baqui Marsischky G., Rolfs A., Richardson A., Kane M., Baqui M. Y., Vannberg F., Weger J., Kramer J., Moreira D., Kel Hu Y., Vanhael J., Hogle C., Jepson D., Williamson J., Zuo D., Raphael J., Hogle C., Jepson A., Kolodner R., ?
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PubMed=9015301; DOI=10.1083/jcb.136.2.299;
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"LMA1 binds to vacuoles at
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MEDLINE=72104597; PubMed=4333602;
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heterodimer of thioredoxin and I(B)2 cooperates
promote yeast vacuole inheritance.";
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with alkyl
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                  687; DOI=10.1074/jbc.274.8.4537;
Labarre J., Toledano M.B.;
hydroperoxide defense propertie
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"A thiol peroxidase is an H2O2 receptor and redox-transducer in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Delaunay A., Isnard A.D., T "H2O2 sensing through oxida EMBO J. 19:5157-5166(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FUNCTION, AND REGULATION PubMed=11013218; DOI=10.1
                                                                                                                                                                                                                    "Yeast thioredoxin. Amino-acid sequence disulfide of thioredoxin I and II.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Thioredoxins are required for protection in the yeast Saccharomyces cerevisiae."; Mol. Microbiol. 46:869-878(2002).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PubMed=12410842;
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                                                                                                                                                                                                                                                                    Hall D.E.,
                                                                                                                                                                                                                                                                                        PubMed=4945270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Role
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                                             Match
                                                                                                   PUNCTION: Participates as a hydrogen donor in rethrough the reversible oxidation of its active disulfide, accompanied by the transfer of 2 election of the involved in many cellular processing the control of the contr
                                                                                        protons. It is involved deoxyribonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  dynamics.
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  100.
ilarity 100.
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ard A.D., Toledano M.B.;
                                                                                                                                                                                                                                                                    A., Holmgren
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                                                                                        synthesis,
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  Score 47; DE Pred. No. 2.4); Mismatches
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                                                                                        repair of oxidatively
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Carroll A.S., Howson R.W
                                                                                                                                                                                                                                                                    Reichard
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Matches 6
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05-JUL-2004
05-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                  TIGRFAMS;
PROSITE; F
STRAIN-ATC 2001 / CBS 138;
PubMed=15229592; DOI=10.1038/nature02579;
Nuion B., Sherman D., Fischer G., Durrens
                                                                            Candida glabrata (Yeast) (Torulopsis glabrata).
Bukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; mitosporic Saccharomycetales; Candida.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PubMed=15001715; DOI=10.1126/science.1095781; Dietrich F.S., Voegeli S., Brachat S., Lerch A., Gate Mohr C., Poehlmann R., Luedi P., Choi S., Wing R.A., Gaffney T.D., Philippsen P.; "The Ashbya gossypii genome as a tool for mapping the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ashbya gossypii (Yeast) (Eremothecium gossypii).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Eremothecium.
MCBI_TaxID=33169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JT 28
20 ASHGO PRELIMINARY;
275CQO;
                                                                Bukaryota; Fungi; Ascomycota;
Saccharomycetales; mitosporic
NCBI_TaxID=5478;
                                                                                                                        OrderedLocusNames=CAGL0K00803g
                                                                                                                                05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Candida glabrata strain_CBS138 chromosome K complete sequence.
                                                                                                                                                                                     )5_CANGA
Q6FND5_CANGA PRELIMINARY;
Q6FND5;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin.like
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; AE016816; AAS51097.1; -; Genomic_DNA. HSSP; P10599; IAUC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Saccharomyces cerevisiae genome.";
Science 304:304-307(2004).
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STRAIN=ATCC 10895;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACL131Wp.
                                          NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Name=ACL131W;
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GO:0005489; F:electron transporter activity; IEA
GO:0006118; P:electron transport; IEA.
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IS; TIGR01068; thioredoxin; 1.

I; PS00194; THIOREDOXIN; 1.
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ilarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                        11396 MW;
                                          [LARGE SCALE GENOMIC DNA]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genome as a tool for mapping the ancient
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Last annotation update)
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Pred. No. 2.
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     Durrens P.,
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     Casaregola S.,
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worrard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
RA Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
RA Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
RA Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
RA Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
RA Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
RA Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
RA Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
RA Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
RA Seniou-Meyer M., Zivanovic Y., Bolotin-Fukuhara M., Thierry A.,
A Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
A Wincker P., Souciet J.-L.;
T "Genome evolution"
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RA Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
RA Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
RA Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
RA Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
RA Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
RA Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
RA Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
RA Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
RA Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
RA Micaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
RA Wellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
RA Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
RA Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
RA Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
RA Geniou-Meyer M., Zivanovic Y., Bolotin-Fukuhara M., Thierry A.,
RA Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
RA Wincker P., Souciet J.-L.;
"Genome evolution in yeasts.";
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Best Local S
Matches 6
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GG; GG:0005489; F:electron transporter activit
GO; GO:0006118; p:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-1ike.
InterPro; IPR012336; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q6BJ80;
25-OCT-2004 (TrEMBLrel. 28, Created)
25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
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SEQUENCE
                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
STRAIN=ATCC 36239 / CBS 767;
PubMed=15229592; DOI=10.1038/nature02579;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Fungi; A
Saccharomycetales;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Debaryomyces hansenii (Yeast) (Torulaspora hansenii)
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Sacc
Saccharomycetales; Saccharomycetaceae; Debaryomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OrderedLocusNames=DEHA0G05126g;
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                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=4959;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q6BJ80_DEBHA PRELIMINARY;
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103 AA;
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Pred. No. 2.4;
0; Mismatches
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QSACNI_CANAL
ID QSACNI_CANAL
ID QSACNI_CANAL
ID QSACNI_C
AC QSACNI;
DT 10-MAY-2
DT 10-MAY-1
RA JONES T.
RA JONES T.
RA GOBERTS
RA ROBERTS
RA ROBERTS
RA ROBERTS
RA ROBERTS
RA GO; GO:(
DR GO; GO:(
DR GO; GO;(
DR GO; GO;(
DR GO; GO;(
DR INTERPRO
DR PFEAM; P
DR PFEAM; P
DR PFEAM; P
PAR SOUTEN
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                                                              Query Match
Best Local S
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                       Dungan J., Kuo A., Newport G., Lan C.-Y., Iijima C., Adegbo A Roberte J., Persson K., Donnelly S., Favoreto S., Tzung K.-
A Jones T., Scherer S., Agabian N.;

"Aunotation of the Genome of Candida albicans.";

"Manotation of the Genome of Candida albicans.";

Submitted (APR-2004) to the EMBL/GenBank/DDBJ databases.

EMBL; AACQ01000032; EAL00485.1; -; Genomic DNA.

GO; GO:0005489; F:electron transporter activity; IEA.

R GO; GO:000518; P:electron transport; IEA.

R GO; GO:0006118; P:electron transport; IEA.

R InterPro; IPR006665; Thioredox dom2.

R InterPro; IPR006665; ThioredoxIn.

R InterPro; IPR006746; ThioredoxIn.

InterPro; IPR012335; Thioredoxin-like.

R InterPro; IPR012335; Thioredoxin-fold.

R Pfam; PF00085; Thioredoxin; 1.

PTGPENEMAG: TYTEOTORGE. Thioredoxin.
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
TIGREAMS; TIGR01068; thioredoxin; 1.
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10-MAY-2005 (TrEMBLrel. 30, Last;
10-MAY-2005 (TrEMBLrel. 30, Last
                                                                                                                                                                                                       PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PubMed=15123810; DOI=10.1073/pnas.0401648101; Jones T., Federspiel N.A., Chibana H., Dungan J., Jones T., Newport G., Thorstenson Y.R., Agabian Davis R.W., Scherer S.; The State of Canada albicans."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=SC5314;
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GO:0005489; F:electron transporter acti
GO:0006118; P:electron transport; IEA.
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                                                              6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WCGPCK 34
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   WCGPCK 6
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                                                              Score 47; DB Pred. No. 2.4
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Pred. No. 2.4;
; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1F533CB671C532EC CRC64;
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                                                                                                                                                                                                       4B78100038A53620 CRC64;
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N., Magee
                                                                     Indels
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RESULT 33

QSFLW1 LACAC

ID QSFLW1 LACAC

RESULW1 LACAC

QSFLW1;

AC QSFLW1;

AC QSFLW1;

DT 10-MAY-2005 (TYENBLYel. 30, La

DE CONTROL SEQUENCE.

OX NCBI TAXID=1579;

RN (1)

RN NUCLEOTIDE SEQUENCE.

RY NUCLEOTIDE SEQUENCE.

RY STRAIN=NCFW;

RN NUCLEOTIDE SEQUENCE.

RY PUBMEd=15671160; DOI=10.1073/;

RA BUCK B.L., MCAULIFIÉE O., Sout!

RA BUCK B.L., MCAULIFIÉE O., Sout!

RA BUCK B.L., MCAULIFIÉE O., Sout!

RA Lick S., Hamrick A., Cano R.,

RT Lactobacillus acidophilus NCFF

RI Tactobacillus ACFG

RMBL; CP000033; AAV42313.1; -
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Q7P4W8 FUSNV PRELIMINARY;
ID Q7P4W8, FUSNV PRELIMINARY;
AC Q7P4W8;
DT 01-MAR-2004 (TrEMBLrel. 26,
DT 01-MAR-2004 (TrEMBLrel. 26,
DT 01-MAR-2004 (TrEMBLrel. 26,
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Best Local S
Matches 6
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Karpatral V., Ivanova N., Anderson I., Reznik G., Bhatta-Gardner W.L., Mikhailova N., Larsen N., D'Souza M., Walu-Gardner W.L., Mikhailova N., Larsen N., D'Souza M., Walu-Gardner R., Overbeek R., Kyrpides N.;
Submitted (JAN-2003) to the EMBL/GenBank/DDBJ databases.
-I- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry w
                        PubMed=15671160; DOI=10.1073/pnas.0409188102; Altermann E., Russell W.M., Azcarate-Peril M.A., B. Buck B.L., McAuliffe O., Souther N., Dobson A., Du Lick S., Hamrick A., Cano R., Klaenhammer T.R.; "Complete genome sequence of the probiotic lactic Lactobacillus acidophilus NCFM."; Proc. Natl. Acad. Sci. U.S.A. 102:3906-3912(2005).
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InterPro; IPR005746; Thioredoxin.
InterPro; IPR006663; Thioredox_dom2
Pfam; PF00085; Thioredoxin; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=ATCC 49256;
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Bacteria; Fusobacteria;
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                                                                                                                                                                                                                                                           Lactobacillus acidophilus.
Racteria; Firmicutes; Lactobacillales;
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        Natl. Acad. Sci. U.S.A. 102:3906-3912(2005). CP000033; AAV42313.1; -; Genomic_DNA.
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103 AA; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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11369 MW; 44
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Fusobacterales; Fusobacteriaceae;
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Pred. No. 2.4;
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4437B8AAD056D7C0 CRC64;
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                                                                                                                         Barrangou
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., Walunas T.,
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                                                                    acid bacterium
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RESULT ORRENGE OF THE CONTROL OF THE
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Best Local S
Matches 6
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2 EMBL; AE009951; AAL94306.1; -; Genomic_DNA.

2 HSSP; P00274; 1T7P.

3 GO; GO:0005489; F:electron transporter activity; IEA.

3 GO; GO:0006118; P:electron transport; IEA.

4 R InterPro; IPR006662; Thioredox dom2.

5 R InterPro; IPR006663; Thioredoxin.

6 R InterPro; IPR006663; Thioredoxin.

7 InterPro; IPR012335; Thioredoxin-like.

8 R InterPro; IPR012335; Thioredoxin-fold.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       QBRH23; FUSNN PRELIMINARY; QBRH23; O1-JUN-2002 (TrEMBLrel. 21, C O1-JUN-2002 (TrEMBLrel. 21, L O1-MAR-2004 (TrEMBLrel. 26, L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE;
Complete
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kapatral V., Anderson I., Ivanova N., Reznik G., Los T., Lykidis i Bhattacharyya A., Bartman A., Gardner W., Grechkin G., Zhu L., Vasieva O., Chu L., Kogan Y., Chaga O., Goltsman E., Bernal A., Larsen N., D'Souza M., Walunas T., Pusch G., Haselkorn R., Fonsten M., Kyrpides N.C., Overbeek R.; "Genome sequence and analysis of the oral bacterium Fusobacterium strain ATCC 25586.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=ATCC 25586;
MEDLINE=21886394; PubMed=11889109;
DOI=10.1128/JB.184.7.2005-2018.2002;
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GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
TIGREPAMS; TIGR01068; thioredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OrderedLocusNames=FN0093;
                                                                                                                                                                                                                                                                        Pfam; PF00085; Thioredoxin; 1. PRINTS; PR00421; THIOREDOXIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fusobacterium nucleatum (subsp. nucleatum).
Bacteria; Fusobacteria; Fusobacterales; Fusobacteriaceae;
                                                                                                                                                                                                                                                TIGRFAMs; TIGR01068; thioredoxin; 1.
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103 AA;
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                                                                                                 Conservative
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                                                                                               Score 47; DB 2
Pred. No. 2.4;
0; Mismatches
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Pred. No. 2.4
0; Mismatches
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                                                                                                                                            DB 2;
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Q74KU7; LACJO
ID Q77KU7;
AC Q74KU7;
AC Q74KU7;
AC Q74KU7;
AC Q74KU7;
DT 05-UUL-2004 (TYEMBLEEL 27,
DE TAIOTED SEQUENCE [LARGE RED 10, 107]
RAM (I)
RAM NUCLEOTIDE SEQUENCE [LARGE RC STRAIN=NCC 533;
RN (I)
RAM PRIDMORE 14983040; DOI=10.107
RA PRIDMORE 14983040; DOI=10.107
RA PRIDMORE 14. C., Zwahlen M.-C.,
RA PITTEE A.-C., Zwahlen M.-C.,
RA PITTEE A.-C., Zwahlen M.-C.,
RAMOLLET B., METCENIE R., KI.
RT "The genome sequence of the
RT Lactobacillus johnsonii NCC
RL POCO. Natl. Acad. Sci. U.S..
DR EMBL; AE017201; AAS08472.1;
DR HSSE; P10599; LAUC.
DR GO; GO:0005118; P:electron
DR GO; GO:0005118; P:electron
DR GO; GO:000518; P:electron
DR GO; GO:000518; P:electron
DR GO; GO:000518; P:electron
DR GO; GO:000518; P:electron
DR InterPro; IPR005663; Thiore
DR InterPro; IPR012335; Thiore
DR PINTS; PR00421; THIOREDOX;
DR PRINTS; PR00421; THIOREDOX;
CM Complete proteome.
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      CCCCCPRRRRRRROCOCOGGEDADA
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GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin, fold.
InterPro; IPR012335; Thioredoxin, fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
TIGREAMS; TIGR01068; thioredoxin; 1.
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097680;
16-OCT-2001
16-OCT-2001
13-SEP-2005
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Pridmore R.D., Barger B., Desiere F., Vilanova D., Barretto C.,
Pittet A.-C., Zwahlen M.-C., Rouvet M., Altermann B., Barrangou R.,
Mollet B., Mercenier A., Klaenhammer T., Arigoni F., Schell M.A.;
"The genome sequence of the probiotic intestinal bacterium
Lactobacillus johnsonii NCC 533.";
Proc. Natl. Acad. Sci. US.A. 101:2512-2517(2004).
EMBL; AE017201; AAS08472.1; -; Genomic_DNA.
HSSP, P10599; IAUC.
                                                                                         NUCLEOTIDE SEQUENCE [MRNA]

MEDLINE=20189621; PubMed=10727087;

MEDLINE=20189621; PubMed=10727087;

Terashima H., Gotoh S., Yagi K., Mizoguchi T.;

Terashima H., Gotoh S., Yagi K., Mizoguchi T.;

"CDNA sequence of bovine thioredoxin.";

DNA Seq. 10:331-333(1999).

INA Seq. 10:331-333(1999).

INA Seq. 10:331-333(1999).

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Bos taurus
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Lactobacillus.
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NCBI_TaxID=9913;
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Mammalia; Eutheria;
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SUBCELLULAR LOCATION: Cytoplasmic
                                 and catalyzes dithiol-disulfide similarity).
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(Rel. 40, Last sequence update)
(Rel. 48, Last annotation updat
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100.0%; Pr
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Laurasiatheria; Cetartiodactyla; Ruminantia;
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Pred. No. 2.4
0; Mismatches
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   (By similarity).
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RESULT 37
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28-FEB-2003
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SEQUENCE
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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HSSP; P10599; 1ERT
SMR; 097680; 1-104
                                                                                               NUCLEOTIDE SEQUENCE [MRNA].

MEDLINE=21576430; PubMed=11719593; DOI=10.1093/molehr/7.12.1159;
Lopata A., Sibson M.C., Enders A.C., Bloomfield K.L., Gregory M.

Trapani G.D., Perkins A.V., Tonissen K.F., Clarke F.M.;

"Expression and localization of thioredoxin during early implant in the marmoset monkey.";
                                                                                                                                                                                                                                                                                                  298៦រី3 ;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-fold.
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
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                                                                                                                                                                                                                                            Name=TXN;
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                    and catalyzes dithiol-disulfide exchange r
SUBCELLULAR LOCATION: Cytoplasmic (By simi
SIMILARITY: Belongs to the thioredoxin fam
SIMILARITY: Contains 1 thioredoxin domain.
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SIMILARITY:
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                                            FUNCTION: Participates in various redox reactions the reversible oxidation of its active center dithiol to and catalyzes dithiol-disulfide exchange reactions. SUBCELLULAR LOCATION: Cytoplasmic (By similarity).
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48, Last annotation
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Contributes to redox potential visimilarity).
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Primates; Platyrrhini;
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RESULT 38
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HSSP; P10599; 1E
SMR; Q9BDJ3; 1-1
                                                                             J. Biol. Chem. 263:9607-9611(1700).

-I- FUNCTION: Participates in various redox reactions reversible oxidation of its active center dithiol and catalyzes dithiol-disulfide exchange reactions.
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SEQUENCE
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       This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                         differentially expressed in normal chicken embryo fibroblasts.";
J. Biol. Chem. 263:9607-9611(1988).
                                                                                                                                                                                                                                                    Name=TXN;
Gallus gallus (Chicken).
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13-SEP-2005
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P08629;
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InterPro;
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ACT_SITE
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TIGRFAMS; TIGR01068; thioredoxin;
                                                                                                                                                                 Jones S.W., Luk K.-C.
                                                                                                                                                                                                                                          Eukaryota; Metazoa;
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IPR006663; Thioredox dom2.
IPR005746; Thioredoxin.
IPR012336; Thioredoxin-like.
IPR012335; Thioredoxin-fold.
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08, Last sequence 48, Last annotation
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Neognathae; Galliformes; Phasianidae; Phasiani
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Deprotonates C-terminal active (By similarity).
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                EMBL; AB022431; BAA37154.1;
HSSP; P10599; 3TRX.
SMR; O97508; 1-104.
                                                               This Swiss-Prot entry is copyright. It is produced through a collaboratio between the Swiss Institute of Bioinformatics and the EMBL outstagion the European Bioinformatics Institute. There are no restrictions on the use as long as its content is in no way modified and this statement is no
                                                                                                                                                                                                                                      Bukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Laurasiatheria; Perissodactyla;
                                                                                                                                                                                                                                                          Equus caballus (Horse)
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PROSITE; PS00194; THIOREDOXIN: 1.
Electron transport; Redox-active center;
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InterPro; IPR00663; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
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SMR; P08629; 1-104.
Ensembl; ENSGALG00000015704;
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        InterPro;
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Thioredox_dom2
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Nucleophile (By similarity).

Nucleophile (By similarity).

Deprotonates C-terminal active s:
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Contributes to redox potential v.
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INIT MET
ACT SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=89008454; PubMed=3170595; Wollman E.E., D'Auriol L., Rimsky L., Wingfield P., Graber P., Dessarps F.; "Cloning and expression of a cDNA for J. Biol. Chem. 263:15506-15512(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                THIO_HUMAN STANDARD; PRT; 104 AA. P10599; Q96KI3; 01-JUL-1989 (Rel. 11, Created) 01-DEC-1992 (Rel. 24, Last sequence update) 13-SEP-2005 (Rel. 48, Last annotation update) Thioredoxin (ATL-derived factor) (ADF) (Surface)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SITE
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InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
                                                                                TISSUE=Lens;
Reddy P.G.,
                                                       Submitted (JUN-2000)
                                                                                                                                                                                                                                                         "ATL-derived factor (ADF), an IL-2 rethioredoxin; possible involvement of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gene 102:221-228(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=91340156; PubMed=187444: Tonissen K.F., Wells J.R.E.; "Isolation and characterization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Name=TXN; Synonyms=TRDX, TRX, TRX1;
Homo sapiens (Human)
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NUCLEOTIDE SEQUENCE
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TIGR01068; thioredoxin; 1.
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104
                                                                                                                                          SEQUENCE
                                                                                Bhuyan D.K.,
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Nucleophile (By similarity).

Nucleophile (By similarity).

Nucleophile (By similarity).

Deprotonates C-terminal active site Cys

(By similarity).
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Contributes to redox potential
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2.5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             thioredoxin-encoding genes.";
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Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
Ak Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
Ak Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
Ak Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
Ak Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
Ak Altechenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Ak Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Schetz T.E.,
Ak Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Schetz T.E.,
Ak Roberts S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Ak Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
Ak Hichards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Ak Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Ak Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
Ak Hiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Ak Rodriguez A.C., Grimwood J., Schwutz J., Myers R.M.,
Ak Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
Ak Rodriguez A.C., Grimwood J., Schmutz J.M., Marra M.A.;
Ak Rodriguez A.C., Grimwood J., Schmutz J.M., Marra M.A.;
Ak Schiernch A., Schein J.E., Jones S.J.M., Marra M.A.;
Brock S.J. M., Marra M.A.;
Brock J. M. J. Jones S.J.M., Marra M.A.;
Ak Schiernch A., Schein J.E., Jones S.J.M., Marra M.A.;
Brock J. Jones S.J.M., Jones S.J.M., Marra M.A.;
Brock J. Jones S.J.M., Jones
oxidized and reduced states of human thioredoxin Structure 2:503-522(1994).
                                              Qin J., Clore ...
"The high-resolution and reduced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE (GENOMIC DNA):
Rieder M.J., Livingston R.J., Daniels M.R., Montoya M.A., Chung M.-W.,
Rieder M.J., Livingston R.J., Daniels M.R., Montoya M.A., Chung M.-W.,
Miyamoto K.E., Mguyen C.P., Nguyen D.A., Poel C.L., Robertson P.D.,
Schackwitz W.S., Sherwood J.K., Witrak L.A., Nickerson D.A.;
"NIEHS SNPs, environmental genome project, NIEHS ES15478, Department
of Genome Sciences, Seattle, WA (URL: http://egp.gs.washington.edu).";
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                Forman-Kay J.D., Clore G.M., Dricoll P.C., Wingfield P., Richards F.M., Gronenborn A.M.; Richards F.M., Gronenborn A.M.; "A proton nuclear magnetic resonance assignment and secondary structure determination of recombinant human thioredoxin.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted
                                                                                                                           STRUCTURE BY NMR.
MEDLINE=95006318; PubMed=7922028;
                                                                                                                                                                                                  Forman-Kay J.D., Clore G.M., Wingfield P., Gronenborn A.M.; "High-resolution three-dimensional structure of reduced recomment thioredoxin in solution."; Biochemistry 30:2685-2698(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=90057393; PubMed=2684271; Forman-Kay J.D., Clore G.M., Dric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=22388257; PubMed=12477932;
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                                                                                                                                                                                                                                                                                                    MEDLINE=91159399; PubMed=2001356;
                                                                                                                                                                                                                                                                                                                           STRUCTURE BY NMR.
                                                                                                                                                                                                                                                                                                                                                                            Biochemistry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRUCTURE BY NMR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         plasma membranes.";
Biochem. Biophys. Res.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Identification of a thioredoxin-related protein associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Martin H., Dean
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PROTEIN SEQUENCE OF 1-14.
MEDLINE=91151337; PubMed=
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and sequencing
d (OCT-2000) to
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                                                                                                                                                                                                                                                                                                                                                                            28:7088-7097(1989).
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                                                     thioredoxin."
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EMBL; X77584; CAA54687.1; -; mRNA.

EMBL; X54539; CAA38410.1; -; Genomic_DNA.

EMBL; X54540; CAA38410.1; JOINED; Genomic_DNA

EMBL; X54541; CAA38410.1; JOINED; Genomic_DNA

EMBL; J04026; AA374596.1; -; mRNA.

EMBL; AV004872; AAF87085.1; -; mRNA.

EMBL; AY004872; AAF87085.1; -; mRNA.

EMBL; AY004872; AAF87085.1; -; mRNA.

EMBL; AF548001; AAG34699.1; -; mRNA.

EMBL; AF548001; AAM33187.1; -; GENOMIC_DNA.

EMBL; BC003377; AAH03377.1; -; GRNA.

EMBL; BC054866; AAH54866.1; -; mRNA.
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MEDLINE=96347359;
Qin J., Clore G.M.
"The solution stru
from Ref-1 reveals
    # FIR ; # FIR 
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MEDLINE=98039128; PubMed=9369469; DOI=10.1021/bi9710048;
Andersen J.F., Sanders D.A., Gasdaska J.R., Weichsel A.,
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Jacquot J.-P., de
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Weichsel A., Gasdaska J.R., Powis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     between
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIINE=96399719; PubMed=8805557; DOI=10.1016/S0969-2126(96)00079-2; Weichsel A., Gasdaska J.R., Powis G., Montfort W.R.; "Crystal structures of reduced, oxidized, and mutated human "Crystal structures of regulatory homodimer.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FUNCTION: Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulfi and catalyzes dithiol-disulfide exchange reactions.

FUNCTION: ADF augments the expression of the interleukin-2 receptor TAC (ILZR/P55).

SUBUNIT: Homodimer.

INTERACTION: Q92905:COP55; NDEXP=5; IntAct=EBI-594644, EBI-594661; SUBCELLULAR LOCATION: Cytoplasmic.

SIMILARITY: Belongs to the thioredoxin family.

SIMILARITY: Contains 1 thioredoxin domain.
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                    1ERU;
1ERW;
1ERW;
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Biophys. Res. Commun. 173:1375-1381(1990)
. F.
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NMR; A=1-104.
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Miginiac-Maslow M.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         way modified and
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PDB; 3TRX; NMR; @=1-104.

PDB; 4TRX; NMR; @=1-104.

IntAct; P10599; -.

SWISS-2DPAGE; P10599; HUMAN.

SWISS-2DPAGE; P10599; -.

PHCI-2DPAGE; P10599; -.

Siena-2DPAGE; P10599; -.

Siena-2DPAGE; P10599; -.

Ensembl; ENGG00000136810; Homo sa
HGNC; HGNC:12435; TXN
H-INVDB; HIX0008875; -.
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01-APR-1993
13-SEP-2005
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"Thioredoxin gene expression is transcriptionally up-regulated by retinol in monkey conducting airway epithelial cells.";

Biochem. Biophys. Res. Commun. 183:170-175(1992).

-!- FUNCTION: Participates in various redox reactions through the reversible exidation of its active center dithiol to a disulfide and catalyzes dithiol-disulfide exchange reactions.

-!- SUNCELLULAR LOCATION: Cytoplasmic (By similarity).

-!- SIMILARITY: Belongs to the thioredoxin family.
Electron
INIT MET
ACT_SITE
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HSSP; P10599; 1ERT.
SMR; P29451; 1-104.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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PIR; JS0667; JS0667.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This Swiss-Prot entry is copyright. It is produced through a celebrate the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict use as long as its content is in no way modified and this statuse as long as its content is in no way modified and this statuse as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status.
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                                                                                                           PRINTS; PR00421; THIOREDOXIN.
TIGRPAMS; TIGR01068; thioredoxin; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
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MEDLINE=92181438; PubMed=1543487;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
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Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,

Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,

A yagi K., Tomaru Y., Hasegawa Y., Nogami A., Schonbach C., Gojobori T.,

Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,

Schriml L.M., Kanapin A., Matsuda H., Batalov S., Beisel K.W.,

Schriml L.M., Kanapin A., Matsuda H., Batalov S., Beisel K.W.,

Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,

A Blake J.A., Bradt D., Brusic V., Chothia C., Gorbani L.E., Cousins S.,

Casting T.A., Fletcher C.F., Forrest A., Frazer K.S.,

A Dalla E., Dragani T.A., Fletcher C.F., Godzik A., Gough J.,

Gassterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,

Gassterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,

Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,

Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,

Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,

Nagashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G.,

Petrovsky N., Pillai R., Pontius J.U., Qi D., Ramachandran S.,
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=129/Sv; TISSUE-Liver;
MEDLINE=95137382; PubMed=7835695; DOI=10.1016/0378-1119(94)00707-Y;
Matsui M., Taniguchi Y., Hirota K., Taketo M., Yodoi J.;
"Structure of the mouse thioredoxin-encoding gene and its processed pseudogene.";
Gene 152:165-171(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE [MRNA].

MEDLINE=89251607; PubMed=2785919;

Tagaya Y., Maeda Y., Mitsui A., Kndo N., Matsui H., Hamuro J.,

Brown N., Arai K.-I., Yokota T., Wakasugi H., Yodoi J.;

Brown N., Arai K.-I., Yokota T., Wakasugi H., Yodoi J.;

"ATL-derived factor (ADP), an II-2 receptor/Tac inducer homologous thioredoxin; possible involvement of dithiol-reduction in the II-2 receptor induction.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       THIO_MOUSE STANDARD; PRT; 104 AA. P10639; Q9DBR0; 01-JUL-1989 (Rel. 11, Created) 01-JUN-1994 (Rel. 29, Last sequence update) 13-SEP-2005 (Rel. 48, Last annotation update) Thioredoxin (ATL-derived factor) (ADF).
                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].
STRAIN=C57BL/6J; TISSUE=Pancreas;
MEDLINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ERRATUM, AND SEQUENCE REVISION.

MEDILINE=94244626; PubMed=8187776;

Tagaya Y., Maeda Y., Mitsui A., Kndo N., Matsui H.,

Brown N., Arai K.-I., Yokota T., Wakasugi H., Yodoi

EMBO J. 13:2244-2244(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniat;
Mammalia; Eutheria; Euarchontoglires;
Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Name=Txn; Synonyms=Txn1; Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SITE
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Redox-active (By similarity).
; C804D5152F8870EB CRC64;
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RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Cassavant T.L., Scheetz T.E.,
RA Brownstein M.J., Ugdin T.B., Toshyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Ugdin T.B., Toshyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Robak S.A., WcEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Sarchan A., Rodriques S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Mhiting M., Madan A., Worng A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Schein J.E., Jones S.J.M., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human rad mouse cDNA sequences";
RI Proc. Natl Road. Sci. U.S.A. 99:16899-16903 (2002).

1 - FUNCTION: Participates in various redox reactions through the contestions of its active center dithiol to a disulfide contestion of its active center dithiol to a disulfide
                                                                                                                                                                                                               EMBL; X77585; CAA54688.1; -; mRNA.

EMBL; D21859; BAA04881.1; -; Genomic DNA.

EMBL; AK007537; BAB25096.1; -; mRNA.

EMBL; AK007790; BAB25256.1; -; mRNA.

EMBL; BC010756; AAH10756.1; -; mRNA.

PIR; JC4068; S04107.

HSSP; P10599; 1ERT.

SMR; P10639; 1-102.

SWISS-2DPAGE; P10639; MOUSE.

Ensembl; ENSMUSG00000050612; Mus musculus.
                 MGI; MGI:98874; Txn1.
GO; GO:0005739; C:mitochondrion; IDA.
InterPro; IPR006662; Thiored.
IntexPro; IPR006663; Thioredox dom2.
IntexPro; IPR012336; Thioredoxin-like.
IntexPro; IPR012335; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TISSUE=Colon;
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as long a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SUBCELLULAR LOCATION: Cytoplasmic. SIMILARITY: Belongs to the thioredo SIMILARITY: Contains 1 thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and catalyzes dithiol-disulfide exchange reactions. FUNCTION: ADF augments the expression of the interleukin-2 receptor TAC (IL2R/P55).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        full-length cDNAs.";
420:563-573(2002).
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THIOREDOXIN; 1.
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Thioredox dom2.
Thioredoxin-like
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RESULT 43
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                   PRINTS; PR00421; THIOREDOXIN.
PROSITE; PS00194; THIOREDOXIN;
Electron transport; Redox-acti
INIT MET 0 0 N
ACT_SITE 31 34 N
SITE 25 25 D
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28-FEB-2003
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CONFLICT
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                                                                                                                                   InterPro;
InterPro;
                                                                                                                                                        EMBL; AF321769; AAK
HSSP; P10599; 3TRX.
                                                                                                                                                                                                                                                     similarity).
-!- SIMILARITY: Belongs to the thioredoxin family.
-!- SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                                                                                                                                                                                          protein.";
Submitted (NOV-2000)
                                                                                                                                                                                                                                                                                                                                               TISSUE=Venom gland;
Lee W., Liu H., Zhang Y.;
"cDNA sequence of Ophiophagus hannah venom gland thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Name=TXN;
Ophiophagus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Thioredoxin
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                                                                                                                                                                                                                          between
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                                                                                                             InterPro;
                                                                                                                       InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                 lapidae; Elapinae;
                                                                                                                                                                                                                                                                                                   FUNCTION: Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulf:
                                                                                                                                                                                                                                                                                          and catalyzes dithiol-disulfide exchange reactions (By
                                                                                                                                                                                                             Swiss-Prot entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the EN European Bioinformatics Institute. There are no resti
                                                                                                 PF00085; Thioredoxin;
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                                                                            PS00194; THIOREDOXIN; 1.
                                                                                                        IPR006662; Thiored.
IPR006663; Thioredox dom2.
IPR012336; Thioredoxin-like.
IPR012335; Thioredoxin_fold.
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Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
a; Squamata; Scleroglossa; Serpentes; Colubroidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
llarity 100.0%;
Conservative 0;
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Pred. No. 2.5;
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S -> C (in Ref. 4; BAB25256)
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Contributes to redox potential
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Contributes to redox potential value
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Deprotonates C-terminal active
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Nucleophile (By similarity).
Deprotonates C-terminal active
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RESULT 44
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; ThioredoxIn.
InterPro; IPR012336; ThioredoxIn-like.
InterPro; IPR012335; ThioredoxIn_fold.
Pfam; PF00085; ThioredoxIn; 1.
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HSSP; P10599; 1F
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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16-OCT-2001 (Rel. 40, Created)
28-FEB-2003 (Rel. 41, Last seq
13-SEP-2005 (Rel. 48, Last ann
                                                                                                                                         PRINTS; PR00421; THICTIGREAMS; TIGRO1068; PROSITE; PS00194; TH
                                                                                                                                                                                                                                                                                                                       SMR; P82460; 1-104
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Yu G.W., Xu J.Y.,
"The cloning and
                                                                                                                                                                                                                                                                                                                                                                                                    removed.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FUNCTION: Participates in various redox reactions to reversible oxidation of its active center dithiol than and catalyzes dithiol-disulfide exchange reactions. SUBCELLULAR LOCATION: Cytoplasmic (By similarity). TISSUE SPECIFICITY: Erythrocytes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SIMILARITY: Belongs to the thioredoxin family. SIMILARITY: Contains 1 thioredoxin domain.
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itted (MAY-2001)
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to Swiss-Prot.
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Nucleophile ()
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leophile (By similarity)
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similarity).
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RESULT 45
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Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam_PF00085; Thioredoxin; 1.
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                                                                                                                                                                                                                                                          reversible oxidation of its active center dithiol t
and catalyzes dithiol-disulfide exchange reactions.
SUBCELLULAR LOCATION: Cytoplasmic.
SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
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P08628; 1-104.
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Matches 6
                 GO; GO:0004791; F:thioredoxin-disulfide re
GO; GO:0006118; P:electron transport; TAS:
InterPro; IPR006662; Thioredox dom2.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
PRINTS; PR00421; THIOREDOXIN.
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE [MRNA]. Xie Z.H., Liu C.Z., He Y.H. Submitted (OCT-2000) to the
                                                                                                                                                                                                                                                                                                                                                                                                                           This Swiss-Prot entry is copyright. It is produced through the European Bioinformatics and the the European Bioinformatics Institute. There are no re-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NIH - Mammalian Gene
Submitted (SEP-2003)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodent
Muroidea; Muridae; Murinae; Rattus.
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SMR; P11232; 1-103.
                                                                                                                                                                                                                                                                               EMMBL, X14478; CAA33019.1; -; mRNA.
EMMBL, AR311055; AAG49923.1; -; mRN
EMBL; BC058454; AAH58454.1; -; mRN
PIR; S04352; S04352.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and catalyzes dithiol-disulfide exchange reactions similarity).
SIMCELLULAR LOCATION: Cytoplasmic (By similarity).
SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
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InterPro; IPR012335; Thioredoxin_fold.
Pfam; PP00085; Thioredoxin; 1.
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SMR; I
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Mammalia; Eutheria; Lauras
Pecora; Bovidae; Caprinae;
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                                                                                                                                                                                                                                                                                                                       MEDLINE=95078463;
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FUNCTION: Participates in various redox reactions reversible oxidation of its active center dithiol and catalyzes dithiol-disulfide exchange reactions
                                                                                                                                                                                                                          similarity).
SUBCELLULAR LOCATION: Cytoplasmic (By simi SIMILARITY: Belongs to the thioredoxin fam SIMILARITY: Contains 1 thioredoxin domain.
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Contributes to redox potential
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Pred. No.
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heria; Cetartiodactyla; Ruminantia;
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MEDLINS-21311952; PubMed=11418146; DDD=10.1016/S00140-6736(00)04403-MEDLINS-21311952; PubMed=11418146; DDD=10.1016/S00140-6736(00)04403-MEDLINS-2131952; PubMed=11. Baba T., Yuzawa H., Kobayashi I., Cui L., Oguchi A., Aoki K.-I., Nagai Y., Lian J.-Q., Ito T., Cui L., Oguchi A., Aoki K.-I., Nagai Y., Lian J.-Q., Ito T., Canamori M., Matsumaru H., Maruyama A., Murakami H., Hosoyama A., Mizutani-Ui Y., Takhhashi N.K., Sawano T., Inoue R.-I., Kaito C., Sekimizu K., Hirakawa H., Kuhara S., Goto S., Yabuzaki J., Kanehisa M., Yamashita A., Oshima K., Furuya K., Yoshino C., Shiba Hattori M., Ogasawara N., Hayashi H., Hiramatsu K., Hunda M., Ogasawara N., Hayashi H., Hiramatsu K., "Whole genome sequencing of meticillin-resistant Staphylococcus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This Swiss-Prot entry is copyright. It is produced through a collaboratic between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use as long as its content is in no way modified and this statement is no
                                                                                                                                                                                                                                                             InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.-like.
InterPro; IPR012336; Thioredoxin-fold.
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Bacteria; Firmicutes; Bacillales; Staphylococcus.
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                                                                                                    proteome; Electron transport; Redox-active center; 29 32 Redox-active (By similarity). 104 AA; 11440 MW; FE61559DB27B3920 CRC64;
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; TIGR01068; thioredoxin; 1.
PS00194; THIOREDOXIN; 1.
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SWISS-DPAGE; P99122; STAAN.
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin, 1.
PRINTS; PR00421; THIOREDOXIN.
TIGRFAM6; TIGR01068; thioredoxin; 1.
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P99122; Q9ZEH4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      aureus.";
Lancet 357:1225-1240(2001).
-!- FUNCTION: Component of the thioredoxin-thioredoxin reductase
                                                                                                                                                                                       This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EW the European Bioinformatics Institute. There are no restruse as long as its content is in no way modified and this s
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Staphylococcus aureus (strain N315).
Bacteria; Firmicutes; Bacillales; St
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-!- SIMILARITY: Belongs to the thioredoxin family.
-!- SIMILARITY: Contains 1 thioredoxin domain.
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RP NUMMEd=15213324; DOI=10.1073/pnas.0402521101;

RA Holden M.T.G., Feil E.J., LindBay J.A., Peacock S.J., Day N.P.J.,

RA Holden M.T.G., Feil E.J., LindBay J.A., Peacock S.J., Day N.P.J.,

RA Enright M.C., Foster T.J., Moore C.E., Hurst L., Atkin R., Barron A.,

RA Enright M.C., Foster T.J., Moore C.E., Hurst L., Atkin R., Barron A.,

RA Churcher C., Clark L., Corton C., Cronin A., Doggett J., Dowd L.,

RA Churcher C., Clark L., Corton C., Cronin A., Doggett J., Dowd L.,

RA Churcher C., Clark L., Corton C., Cronin A., Doggett J., Dowd L.,

RA Churcher C., Harris B., Hauser H., Holroyd S., Jagels K.,

RA James K.D., Lennard N., Line A., Mayes R., Moule S., Mungall K.,

AN James K.D., Lennard N., Line A., Mayes R., Moule S., Mungall K.,

AN Ormond D., Quail M.A., Rabbinowitsch E., Rutherford K.M., Sanders M.,

Sharp S., Simmonds M., Stevens K., Whitehead S., Barrell B.G.,

RA Spratt B.G., Parkhill J.,

"Complete genomes of two clinical Staphylococcus aureus strains:

"T'Complete genomes of two clinical Staphylococcus aureus strains:

"Complete genomes of two clinical Staphylococcus aureus strains:

"Complete genomes of two clinical Staphylococcus aureus strains:

"Complete genomes of two clinical Staphylococcus aureus strains:

"T'Complete genomes of two clinical Staphylococcus aureus strains:

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10-MAY-2005
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                                                                                              Q6GA69;
10-MAY-2005
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10-MAY-2005
                                                                                                                                                                               THIO_S
Staphylococcus aureus Bacteria; Firmicutes; NCBI_TaxID=282459;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; ThioredoxIn.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
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                                                             Name=trxA; OrderedLocusNames=SAS1079
                                                                                Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and catalyzes dithiol-disulfide exchange reactions similarity). SIMILARITY: Belongs to the thioredoxin family. SIMILARITY: Contains 1 thioredoxin domain.
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Bacillales; Stap
                        (strain MSSA476)
Bacillales; Stap
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Pred. No. 2.5;
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PubMed=15213324; DOI=10.1073/pnas.0402521101;

Holden M.T.G., Feil E.J., Lindsay J.A., Peacock S.J., Day N.P.J.

Baright M.C., Foster T.J., Moore C.E., Hurst L., Atkin R., Barro

Bason N., Bentley S.D., Chillingworth C., Chillingworth T.,

Churcher C., Clark L., Corton C., Cronin A., Doggett J., Dowd L.

Feltwell T., Hance Z., Harris B., Hauser H., Holroyd S., Jagels

James K.D., Lennard N., Line A., Mayes R., Moule S., Mungall K.,

Ormond D., Quail M.A., Rabbinowitsch E., Rutherford K.M., Sanderi

Sharp S., Simmonds M. Stevene v. whithers A.
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PROSITE; F
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PRINTS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Complete genomes of two clinical Staphylococcus aureus evidence for the rapid evolution of virulence and drug Proc. Natl. Acad. Sci. U.S.A. 101:9786-9791(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ormond D., Quail M.A., Kaupanora, Whitehead S., Sharp S., Simmonds M., Stevens K., Whitehead S., Spratt B.G., Parkhill J.;
Spratt B.G., Parkhill J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (1)
              Uziel O., Borovok I., Schreiber R., Cohen G., Aharonowit "Transcriptional analysis of the thioredoxin (trxA) and reductase (trxB) genes in Staphylococcus aureus ", Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                       Staphylococcus aureus. Bacteria; Firmicutes;
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                                                                                                                                                                                             NCBI_TaxID=1280;
                                                                                                                                                                                                                                                                         Name=trxA;
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                                                                                                                               STRAIN-Oxford;
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SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
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IPRO06663; Thioredox dom2.
IPRO05746, Thioredoxin.
IPRO12336; Thioredoxin-like.
IPRO12335; Thioredoxin_fold.
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1 (TRX).
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Best Local S
Matches 6
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PP00085; Thioredoxin; 1.
PR.NTS; PR00421; THIOREDOXIN; 1.
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HSSP; P002
InterPro;
InterPro;
EMBL;
HSSP;
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29-MAR-2004 (Rel. 43, Last seq
10-MAY-2005 (Rel. 47, Last ann
Thioredoxin (TRX)
                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
MEDLINE=22040717; PubMed=12044378; DOI=10.1016/S0140-6736(02)01
Baba T., Takeuchi F., Kuroda M., Yuzawa H., Aoki K.-I., Oguchi
Baba T., Takeuchi F., Kuroda M., Kuroda H., Cui L.,
Negai Y., Iwama N., Asano K., Naimi T., Kuroda H., Cui L.,
                                                                                                                                                                                                                                                                                                                                                              Staphylococcus aureus Bacteria; Firmicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use
                                                                                                                                                                                                                                        Yamamoto K., Hiramatsu K.;
"Genome and virulence determinants of high virulence community-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Electron transport;
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                                          removed.
                                                       use as
                                                                                 between
                                                                                                                                                                                                                                      acquired MRSA."
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                                                                                                                     similarity).
SIMILARITY: Belongs t
SIMILARITY: Contains
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SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                                 cet 359:1819-1827(2002). FUNCTION: Component of the thioredoxin-thioredoxin reductase system. Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulf and catalyzes dithiol-disulfide exchange reactions (By
                                                                  Swiss-Prot entry is copyright. It is produ
een the Swiss Institute of Bioinformatics
European Bioinformatics Institute. There a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Swiss-Prot entry is copyright. It is produced through a collaboeen the Swiss Institute of Bioinformatics and the EMBL outsta European Bioinformatics Institute. There are no restrictions o as long as its content is in no way modified and this statement
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              BAB94893
                                                                                            entry is copyright. It is produced through
                                                                                                                        Belongs to the thioredoxin family. Contains 1 thioredoxin domain.
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                                                       content
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Redox-active center; Transport.
32 Redox-active (By similarity)
11440 NW; FE61559DB27B3920 CRC64;
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                                                                                                                                                                                                                                                                                                                                                           (strain MW2).
Bacillales; Staphylococcus.
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annotation updat
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Pred. No.
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              Genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                    GENOMIC DNA).
8; DOI=10.1016/S0140-6736(02)08713-5;
                                                       g
                                                     way modified
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Complete DISULFID SEQUENCE

104 AA;

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proteome; Electron transport; Red
29 32 Redox-active

ansport; Redox-active center; edox-active (By similarity). DBBCE61D2DA7E770 CRC64;

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UR InterPro; IPR006662; Thiored.

DR InterPro; IPR006663; Thioredox dom2.

DR InterPro; IPR005746; Thioredoxin.

DR InterPro; IPR012336; Thioredoxin-fold.

DR Pfam; PF00085; Thioredoxin, fold.

DR PRINTS; PR00421; THIOREDOXIN

R TIGRAMS; TIGR01068; **

PROSTIE; PS0077
RESULT TO THE PROPERTY OF THE 
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STAEP
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DISULFID
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=22832016; PubMed=12950922;
DOI=10.1046/j.1365-2958.2003.03671.x;
DOI=10.1046/j.1365-2958.2003.03671.x;
Zhang Y.-Q., Ren S.-X., Li H.-L., Wang Y.-X., Fu G., Yang J.,
Qin Z.-Q., Miao Y.-G., Wang W.-Y., Chen R.-S., Shen Y., Chen Z.,
Yuan Z.-H., Zhao G.-P., Qu D., Danchin A., Wen Y.-M.;
"Genome-based analysis of virulence genes in a non-biofilm-forming
Staphylococcus epidermidis strain (ATCC 12228).";
Mol. Microbiol. 49:1577-1593(2003).
-I- FUNCTION: Component of the thioredoxin-thioredoxin reductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; ThioredoxIn.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Staphylococcus epidermidis.
Bacteria; Firmicutes; Bacil
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Name=trxA; OrderedLocusNames=SE0838;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  system. Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulfide and catalyzes dithiol-disulfide exchange reactions (By similarity).

SIMILARITY: Belongs to the thioredoxin family.

SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                                                                                                                                                                                                                                                                s Swiss-Prot entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the EN Entropean Bioinformatics Institute. There are no restrant as long as its content is in no way modified and this s
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43, Last sequence
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Pred. No. 2.5;
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RESULT 56
Q6C399 YARLI
ID Q6C399-YARLI PRELIMINARY; PRT;
AC Q6C399;
DT 25-OCT-2004 (TrEMBLrel. 28, Created)
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Best Local S
Matches 6
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Y PubMed=15229592; DOI=10.1038/nature02579;

A Dujon B., Sherman D., Fischer G., Durrens P., Casaregola S.,

A Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,

A Goffard N., Frangeul L., Aigle M., Anthouard V., Babbur A., Barbe V.,

A Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,

A Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,

Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,

A Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,

A Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,

A Micaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,

A Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,

A Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos R.,

A Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,

Zeniou-Meyer M., Zivanovic Y., Bolotin-Fukuhara M., Thierry A.,

Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,

Mincker P. Scholer J.-I.
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Best Local Similarity
Matches 6; Conserv
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EMBL; CR392125; CAG99773.1; -; Genomic DNA.

GO; GO:0005489; F:electron transporter activity; IEF
GO; GO:0006118; P:electron transport; IEA.

InterPro; IPR006662; Thiored.

InterPro; IPR006663; Thioredoxin.

InterPro; IPR005746; Thioredoxin-like.

InterPro; IPR012336; Thioredoxin-like.

InterPro; IPR012336; Thioredoxin-fold.

Pfam; PF00085; Thioredoxin; 1.

PR1NTS; PR0421; THIOREDOXIN.

PR1NTS; PR0421; THIOREDOXIN.
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25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similar to
                                                                                                                                                                                                                                                                                     TIGRFAMS; TIGR01068; thioredox: PROSITE; PS00194; THIOREDOXIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Genome evolution in yeasts.";
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6; Conserv
                                                                                                                                                   1 WCGPCK 6
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Pred. No. 2.5;
0; Mismatches
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Pred. No.
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F., de Daruvar A.,
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RESULT
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Best Local S
Matches 6
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GO; GO:0005489; F:electron transporter activit
GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR006662; Thioredox dom2.
InterPro; IPR006663; Thioredoxin.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
                                                                                                                                                                                                                                                                                                                               Q5KK55_CRYNE
Q5KK55;
10-MAY-2005 (
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SEQUENCE
                                                                                                                                            Cryptococcus neoformans var. neoformans JBC21.
Eukaryota; Fungi; Basidiomycota; Hymenomycetes; Heterobasidiomycetes;
Tremellomycetidae; Tremellales; Tremellaceae; Filobasidiella.
NCBI_TaxID=214684;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=CLIB 122 / E 150;
PubMed=15229592; DOI=10 1038/nature02579;
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Saccharomycetales; Dipodascace
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                                                                                                                                                                                                                                                                 10-MAY-2005
Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bouchier C., Caudron B., S
Wincker P., Souciet J.-L.;
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                                                                                     STRAIN=JEC21;
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                                       Loftus B., Amedeo P.,
Van Aken S., Fraser C
                                                                                                                                                                                                                                                                                                          10-MAY-2005
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(TrEMBLrel. 30, 1
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                                                           Roncaglia P., Vamathevan J., Utterback
                    the EMBL/GenBank/DDBJ databases
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Last annotation update)
2), putative.
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Pred. No. 2.5
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Saccharomycotina;
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QSSW38 C
AC QSSW38;
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DT 13-SEP-2
DT 13-SEP-2
DT 13-SEP-2
DT 13-SEP-2
DT 13-SEP-2
CO ENKARYOUT
OC Tremello
OC Tremello
OX NCBI TAX
RN NUCLEOTI
RC STRAIN-B
RA Wickes B
RT "Cryptoo
RL Submitte
CC Prell
DR GO; GO:0
DR GO; GO:0
DR InterPro
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR01663; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
TIGRPAMS; TIGR01068; thioredoxin; 1.
PROCEETER PROCESSION (1)
                                                                                                                                                                                                                                           STRAIN=8.3501A;
STRAIN=8.3501A;
Fung E., Hyman R.W., Rowley D., Bruno D., Miranod M., Fung E., Hyman R.W., Rowley D., Bruno D., Miranod M., Fung E., Hyman R.W., Pu J., Davis R.W.;
Wickes B.L., Fu J., Davis R.W.;
"Cryptococcus neoformans serotype D sequencing.";
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13-SEP-2005 (
13-SEP-2005 (
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                                                                       InterPro;
InterPro;
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InterPro;
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Bukaryota; Fungi; Basidiomycota;
Tremellomycetidae; Tremellales; "NCBI_TaxID=283643;
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Science 307:1321-1324(2005).
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               preliminary data.

MBL; AAEY0100013; EAL22161.1; -; Genomic O; GO:0005489; F:electron transporter aci O; GO:0006118; F:electron transport; IEA (InterPro; IPR006662; Thiored. InterPro; IPR006663; Thioredox dom2. InterPro; IPR0054663; Thioredoxin. InterPro; IPR00546; Thioredoxin. InterPro; IPR012336; Thioredoxin-like. InterPro; IPR012335; Thioredoxin-fold.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypothetical protein.
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RESULT 60
Q9NPK9 PLAFA
ID Q9NPK9 PLAFA PRELIMINARY;
AC Q9NPK9;
DT 01-OCT-2000 (TrEMBLrel. 15,
DT 01-MAR-2004 (TrEMBLrel. 26,
DT 01-MAR-2004 (TrEMBLrel. 26,
DE Thioredoxin.
GN Name=trx;
GN Plasmodium falciparum.
OC Eukaryota; Alveolata; Apicot
OX NCBI_TaxID=5833;
RN [1]
RP NUCLEOTIDE SEQUENCE.
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PROSITE; I
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GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR005736; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin, 1.
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HSSP; P80028; 1EP7.
SMR; Q9NIR2; 1-103.
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EMBL; AF202664; AAF34541.1; -; mRNA.
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Eukaryota; Alveolata;
NCBI_TaxID=5833;
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Q9NIR2;
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MEDLINE=20568274; Po
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PubMed=11013257; DOI=10.1074/jbc.M007633200; rmer R.H., Turbachova I., Iozef R., Becker K.
                                                                                                                                                                                                                                                                                                                        100.0%;
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                                               Apicomplexa; Haemosporida; Plasmodium
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XL Submitted (MAY-2000) to the EMBL/GenBank/DDBJ databases.

RMBL; AJZ77839; CAB90828.1; -; mRNA.

RGO; GO:00051489; F:electron transporter activity; IEA.

RGO; GO:000518; P:electron transporter activity; IEA.

RGO; GO:000618; P:electron transport; IEA.

RInterPro; IPR006662; Thiored.

R InterPro; IPR006663; Thioredox dom2.

R InterPro; IPR012336; Thioredoxin.

R InterPro; IPR012336; Thioredoxin-like.

R InterPro; IPR012335; Thioredoxin-fold.

R Pfam; PF00085; Thioredoxin; 1.

R Pfam; PF00085; Thioredoxin; 1.

R PRINTS. DB00431. "MIOREDOXIN."
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Best Local :
                      GO; GO:0005489; F:electron transporter a GO; GO:0006118; P:electron transport; IE InterPro; IPR006662; Thiored. InterPro; IPR006663; Thioredox dom2. InterPro; IPR005746; Thioredoxin—InterPro; IPR012336; Thioredoxin—like. InterPro; IPR012335; Thioredoxin—fold. Pfam; PF00085; Thioredoxin; 1. PRINTS; PR00421; THIOREDOXIN; UNKNOWN; ITGRPAMs; TIGR01068; Thioredoxin; 1.
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Q7RH10;
01-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-22255706; PubMed-12368865; DOI=10.1038/nature01099; Carlton J.M., Angiuoli S.V., Suh B.B., Kooij T.W., Pertea M., Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L., Peterson J.D., Pop M., Koosack D.S., Shumway M.F., Bidwell S.L., Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V., Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V., Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M., Florens L., Yates J.R. III, Raine J.D., Sinden R.B., Harris M.A., Cunningham D.A., Preiser P.R., Bergman L.W., Vaidya A.B., van Lin L.H., Janse C.J., Waters A.P., Smith H.O., White O.R., van Lin L.H., Janse C.J., Waters A.P., Smith H.O., White O.R., Salzberg S.L., Venter J.C., Fraser C.M., Hoffman S.L., Gardner M.
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Krnajski Z., Gilberger T.W.,
Submitted (MAY-2000) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Genome sequence and comparative analysis parasite Plasmodium yoelii yoelii."; Nature 419:512-519(2002).
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TIGREPAMS; TIGRO1068; thioredoxin; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
SEQUENCE 104 AA; 11716 MW; 01299
                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; AABL01001248; EAA16007.1; -; Genomic_DNA HSSP; P80028; 1EP7. SMR; Q7RH10; 1-103.
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Eukaryota; Alveolata; Api
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Q6B963;
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Thioredoxin
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Thioredoxin.
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pDB; 1SYR; X-ray; A/B/C/D/E/F/G/H/J/J/K/L=1-104.
GO; GO:0005489; F:electron transporter activity;
GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR006663; Thioredoxin.
InterPro; IPR012336; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-fold.
InterPro; IPR012335; Thioredoxin-fold.
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                                        Name=Trx-2;
Schistosoma
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NCBI_TaxID=36329;
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Eukaryota; Metazoa;
Schistosomatoidea;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nature 419:498-511(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Genome sequence of falciparum.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.
MEDLINE=22255705; Pu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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                                                                                                                                                                                                                   SCHMA
                                                                                                                                                                                                                                                                                                                                                                                                               Local
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                                                                                                                                                                                                                                                                                                                                              1 WCGPCK 6
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mansoni (Blood fluke).
Metazoa; Platyhelminthes;
toidea; Schistosomatidae;
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  Trematoda; Digenea;
Schistosoma.
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S., Paulsen I.T., James K.,
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Best Local S
Matches 6
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"The full-length cDNA sequences of Schistosoma japonicum submitted (NOV-2004) to the EMBL/GenBank/DDBJ databases. Submitted (NOV-2004) to the EMBL/GenBank/DDBJ databases. EMBL; AY815296; AAW27028.1; -; mRNA.

GO; GO:0005489; F:electron transporter activity; IEA.
GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-fold.
Pfam; PP00085; Thioredoxin, 1.
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10-MAY-2005 (TrEMBLrel. 30,
10-MAY-2005 (TrEMBLrel. 30,
10-MAY-2005 (TrEMBLrel. 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hypothetical protein.
Schistosoma japonicum (Blood fluke).
Eukaryota; Metazoa; Platyhelminthes;
Schistosomatoidea; Schistosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; ThioredoxIn.
InterPro; IPR012336; ThioredoxIn-like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
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Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases.
EMBL, AY679828, AAT76629.1; -; mRNA.
GO; GO:0005489; F:electron transporter activity; IEA.
GO; GO:0006118; P:electron transport; IEA.
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TIGREAMS; TIGRO1068; thioredoxin; 1.
PROSTITE; PS00194; THIOREDOXIN; 1.
SEQUENCE 104 AA; 11995 MW; 8D36E
                                                                                                                                                                                                                                                                                                      Hypothetical SEQUENCE 1
                                                                                                                                                                                                                                                                                                                                                TIGRFAMS; TIGR01068; thioredox. PROSITE; PS00194; THIOREDOXIN;
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Beers L.F., Williams
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TIGRFAMs; TIGR01068; thioredoxin; 1.
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Pred. No. 2.5
0; Mismatches
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Pred. No. 2.5;
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                                                                                                                                                                                                     Mismatches
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databases.
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QHNM5;
13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
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Q48985;
01-NOV-1996 (TrEMBLrel. 01,
01-NOV-1996 (TrEMBLrel. 01,
01-MAR-2004 (TrEMBLrel. 26,
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                                preliminary data.
EMBL; AAFJ01000019.
                                                                                                                                                                   Nelson K.E.;
"Major structural and novel potential virulence mechanisms genomes of multiple Campylobacter species.";
Submitted (DEC-2004) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                      Fouts D.E., Mongodin E.F., Mandrell R.E., Miller W.G., Rasko D.A.,
Jacques R.J., Brinkac L.M., DeBoy R.T., Parker C.T., Daugherty S.C.,
Dodson R.J., Durkin A.S., Madupu R.R., Sullivan S.A., Shetty J.U.,
Ayodeji M.A., Shvartsbeyn A.A., Schatz M.C., Badger J.H., Fraser C.M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Campylobacter upsaliensis RM3195.
Bacteria, Proteobacteria; Epsilonproteobacteria; Campylobacterales; Campylobacteraceae; Campylobacter.
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"Exploring the Mycoplasma capricolum genome: a minimal cell reveals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mycoplasma capricolum.
Bacteria; Firmicutes; Mollicutes;
   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name=trx; ORFNames=CUP1008;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thioredoxin
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GO; GO:0006118; P:electron transport; IEA.
GO; GO:0006810; P:transport; IEA.
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Mol. Microbiol. 16:955-967(1995).
EMBL; Z33053; CAA83726.1; -; Genomic_DNA.
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STRAIN=ATCC 27343;
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InterPro; IPR005746; Thioredoxin.
InterPro; IPR006663; Thioredox_dom2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PIR; S77780; S77780.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=96059641; PubMed=7476192;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=2095;
                                                                                                         CAUTION: The sequence shown her 
EMBL/GenBank/DDBJ whole genome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE.
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5 (TrEMBLrel. 31,
5 (TrEMBLrel. 31,
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   EAL52411.1; -; Genomic_DNA.
11271 MW; 3056CAA6990B8F74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C133AFEBF027A001 CRC64;
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Q1-MAR-2004 ('01-MAR-2004 ('01-M
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"NADH-dependent bacterial thioredoxin reductase-like protien, conjunction with a glutaredoxin homologue, form a unique pero (AhpC) reducing system in Clostridium pasteurianum.";
Biochemistry 41:1990-2001(2002)
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01-FEB-2005 (TrEMBLrel.
01-FEB-2005 (TrEMBLrel.
Putative thioredoxin.
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TIGREAMS; 110...
TOCTITE; PS00194;
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Bacteria; Firmicutes; Clo
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                                                                                                                                                      Bacteroidaceae; Ba
NCBI_TaxID=272559;
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Q5LHJ1;
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PRINTS; PR00421; THIOREDOXIN.
TIGRPAMS; TIGR01068; thioredoxin; 1.
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                                             NUCLEOTIDE SEQUENCE
PubMed=15746427; DOI
                                                                                                                                                                                                                                             Bacteroides fragilis (strain ATCC 25285 / NCTC 9343).
Bacteria; Bacteroidetes; Bacteroidetes (class); Bacteroidales;
                                                                                                                                                                                                                                                                                                                            Name=trxA; OrderedLocusNames=BF0642;
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HSSP; P00274; 1F6M.
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    Cerdeno-Tarraga
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BACFN
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        A.-M.,
                                             DOI=10.
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                                             [LARGE SCALE GENOMIC DNA]
[=10.1126/science.1107008;
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        Patrick
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PIR data bank
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Pred. No.
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6A60C767560E7ED2 CRC64;
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        Crossman L.C.,
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RESULT 70

Q81718_BACCR
Q81718; BACCR PRELIMINARY;
AC Q81718;
DT 01-UUN-2003 (TrEMBLrel. 24,
DT 01-UUN-2004 (TrEMBLrel. 24,
DT 01-MAR-2004 (TrEMBLrel. 26,
DE Thioredoxin.
GN OrderedLocusNames=BC4521;
OS Bacillus cereus (strain ATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q4L5F01
ID4L5F01
IQ4L5F01
IQ4L5F01
IQ4L5F01
IQ4
AC Q4

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InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox_dom2.
InterPro; IPR005746; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin, 1.
PRINTS; PR0041; THIOREDOXIN.
TICRPAMS; TICR01068; thioredoxin; 1.
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Q4L5F0;
13-SEP-2005
13-SEP-2005
13-SEP-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Whole genome sequencing of Staphylococcus haemolyticus uncovers extreme plasticity of its genome and dynamism in the evolution of human-colonizing staphylococcal species.";
Submitted (DEC-2003) to the EMBL/GenBank/DDBJ databases.
EMBL, AP006716; BAE05125.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene expression.";
Science 307:1463-1465(2005).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Takeuchi F., Watanabe S., Baba T., Yu:
Morimoto Y., Kuroda M., Takahashi M.,
Lee J.C., Hiramatsu K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Staphylococcus haemoly Bacteria; Firmicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
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                                                                     (TrEMBLrel. 24, (TrEMBLrel. 24, (TrEMBLrel. 26,
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Lales; Staphylococcus
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Pred. No. 2.5
0; Mismatches
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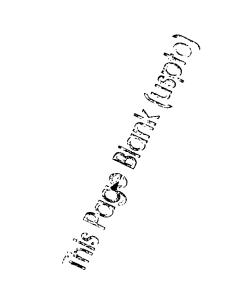
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OC Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus;
OX NCBI_TaxID=226900;
RN [1]
RN NCELCOTIDE SEQUENCE.
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RN NEDLINE=22660415; PubMed=12721630; DOI=10.1038/nature01582;
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RX Agatral V., Bhattacharyya A., Reznik G., Mikhailova W., Lapidus A.,
RX Chu L., Mazur M., Goltsman E., Lareen N., D'Souza M., Malunas T.,
RX Grechkin Y., Pusch G., Haselkorn R., Fonstein M., Ehrlich S.D.,
RX Grechkin Y., Pusch G., Haselkorn R., Fonstein M., Ehrlich S.D.,
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RX Grechkin Y., Pusch G., Haselkorn R., Fonstein M., Ehrlich S.D.,
RX Grechkin Y., Pusch G., Haselkorn R., Fonstein M., Ehrlich S.D.,
RX GO: GO:0005489; Fielectron transporter activity; IEA.
DR GO: GO:0006410, Pitransport; IEA.
DR GO: GO:0006610, Pitransport; I
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Run

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Result
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Sequence 5, Appara-
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; Patent No. 5985261
; Patent No. 5985261
; GENERAL INFORMATION:
; APPLICANT: Dae, Kumuda C.
TITLE OF INVENTION: USE OF THIOREDOXIN-LIKE MOLECULES FOR
TITLE OF INVENTION: INDUCTION OF MASOD TO TREAT OXIDATIVE DAMAGE
; NUMBER OF SEQUENCES: 5
; CORRESPONDENCE ADDRESSE: Sheridan Ross P.C.
ADDRESSEE: Sheridan Ross P.C.
Query Match
Best Local Similarity
6; Conserva
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NAME: Connell, Gary J.
REGISTRATION NUMBER: 32,020
REFERENCE/DOCKET NUMBER: 2879
TELECOMMUNICATION INFORMATION:
TELEPHONE: 303/863-9700
                                                                                                                                                      TELEFAX: 303/863-0223
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: F10ppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/883,804
                                                                                             MOLECULE TYPE:
                                                                                                          TYPE: amino acid TOPOLOGY: linear
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CLASSIFICATION: 514
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              100.0%; Score 47; DB 1; Length 6; 100.0%; Pred. No. 4.6e+05; Live 0; Mismatches 0; Indels
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Sequence 29, Appli
Sequence 6, Appli
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                 Gaps
                 0
            GENERAL INFORMATION:
APPLICANT: Tuite, Michael F.
APPLICANT: Freedman, Robert B
APPLICANT: Markus, Henry Z.
APPLICANT: Schultz, Loren D.
APPLICANT: Montgomery, Donna I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 2
US-08-627-907A-6
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                                                                                                                                         US-07-901-713A-3
                                                                                                                                                        RESULT 3
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                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                     Best
                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: LL.
STREET: New York
CITY: New York
TISA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 6 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U
                                                                                                                                                                                                                                                                                 Local
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                                                                                                                                                                                                                                                                   Similarity 6; Conserv
                 Montgomery, Donna L.
                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                               not relevant
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Best Local Similarity
Matches 6; Conser
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Patent No. 6
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                                                                                                                                                                                                                      SOPTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/750,142B
FILING DATE: 25-NOV-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/SE96/00365
FILING DATE: 22-MAR-1996
APPLICATION NUMBER: SE9501067-4
FILING DATE: 24-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/07/901,713A
CURRENT FILING DATE: 1992-06-12
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                        NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 0627
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
TELEPAX: 617-542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: METHOD FOR INCREASING PRODUCTION OF TITLE OF INVENTION: DISULFIDE BONDED RECOMBINANT PROTEINS BY SACCAROMYCES TITLE OF INVENTION: CEREVISIAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 18469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatibl
                                                                                              TELEFAX: 617-542-8906 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
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ADDRESSEE: Fish & Richardson, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Sarnstrand, Bengt
TITLE OF INVENTION: NEW PEPTIDES WITH
TITLE OF INVENTION: IMMUNOMODULATORY EFFECTS
                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 225 I
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                   TOPOLOGY:
                                   amino acid
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Briksson, Tomas
Lindvall, Magnus
Sarnstrand, Bengt
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3YSTEM: Windows95
peptide
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Pred. No. 4.6e+05;
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5210073-1
;Patent No. 5210073
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TITLE OF INVENTION: METHOD FOR TRREATING CANCER THERAPY; RADIATION DAMAGE OR ARTERIOSCLEROSIS USING HUMAN ADF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO:1:
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Best Local Similarity
Matches 6; Conserv
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Matches 6; Conserv
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                                                                                                                                                                      OPERATING SYSTEM: DOS
SOFTWARE: FRATSEQ for Windows Version 2.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/826,910
FILING DATE: Herewith
CLASSIFICATION = 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
APPLICANT: Shah, Purvi
                                                    NAME: Billings, Lucy J.

NAME: Billings, Lucy J.

REGISTRATION NUMBER: 97-0

RELECOMMUNICATION INFORMATION:

TELEPHONE: 415-855-0555

TELEPAX: 415-845-4166
                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/589,616
SEQUENCE CHARACTERISTICS:
LENGTH: 105 amino acids
                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                  ZIP:
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                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Palo Alto
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ilarity 100.0%;
Conservative 0
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Pred. No.
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Pred. No. 0.75;
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                                                                                                          US-08-826-910-4
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                                                       Matches
                                                                    Best Local Similarity
                                                                               Query Match
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APPLICANT: Hillman, Je
APPLICANT: Shah, Purvi
                                                                                                                                                                                                                                                                      CLASSIFICATION: 530
PRIOR APPLICATION UNDER:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                 TELEX:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/826,910
FILING DATE: Herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: THIOREDOXIN-LIKE PROTEIN NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc. STREET: 3174 Porter Drive
                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 105 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
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CITY: Palo Alto
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                                                                                                                                                                                                                                                             TELEFAX: 415-845-4166
                                                                                                                                                                                          TYPE: amino acid
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                            WCGPCK 6
 WCGPCK 36
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                                                                 Score 47; DB 1; Length 105; Pred. No. 4.4;
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                                                      Mismatches
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RESULT 8

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NUMBER OF SEQ ID NOS: 74
SOFTWARE: PatentIn Ver. 2.
SEQ ID NO 68
LENGTH: 105
TYPE: PRT
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                                                                                                                             RESULT 10
US-08-386-729A-10
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                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
***ches 6; Conserva
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; ORGANISM: Homo sapiens
US-09-513-999C-8037
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US-09-601-144-68
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                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent NO. 6783961
FILE REFERENCE: 59 US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 8037
LENGTH: 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Wright, Jim A.
APPLICANT: Wright, Jim A.
APPLICANT: Young, Aiping H.
APPLICANT: Lee, Yoon S.
TITLE OF INVENTION: OLIGONUCLECTIDE SEQUENCES COMPLEMENTARY TO THIOREDOXIN
TITLE OF INVENTION: AND THIOREDOXIN REDUCTASE GENES AND METHODS OF USING
TITLE OF INVENTION: SAME TO MODULATE CELL GROWTH
FILE REFERENCE: 683-112US-A
CURRENT APPLICATION NUMBER: US/09/601,144
CURRENT FILING DATE: 2000-10-18
CURRENT FILING DATE: 2000-10-18
                                                                    Sequence 10, Application US/08386729A Patent No. 5753435 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -09-00-
Sequence 68, App...
Sequence 68, App...
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APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/073,196
PRIOR FILING DATE: 1998-01-30
                                    APPLICANT:
   APPLICANT:
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mes 6; Conserv
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Cohen, Gerald
Bovenberg, Roelof A. L.
                    Aharonowitz, Yair
Van Der Voort, Lucia H.
Cohen, Gerald
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Pred. No.
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RESULT 11
US-07-745-382-22
J Sequence 22, Application US/07745382
J Patent No. 5270181
J Patent T THOMARTON:
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Best Local Similarity
Whiches 6; Conserve
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APPLICATION NUMBER: US 08/274,043
FILING DATE: 12-JULY-1994
APPLICATION NUMBER: US 07/820,688
FILING DATE: 24-MARCH-1992
APPLICATION NUMBER: PCT/NL91/000101
FILING DATE: 18-JUNE-1991
APPLICATION NUMBER: EP 90201598.1
FILING DATE: 18-JUNE-1990
ATTORNST/AGENT INFORMATION:
NAME: Rae-Venter, Barbara
REGISTRATION NUMBER: 32,750
REFERENCE/DOCKET NUMBER: GBRO-024/02US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Nan, Helen
APPLICANT: Kattevilde
APPLICANT: Pallissa,
TITLE OF INVENTION: A
TITLE OF INVENTION: C
                   APPLICANT: McCoy, John
APPLICANT: LaVallie, Edward
TITLE OF INVENTION: Peptide and Protein Fusions To
TITLE OF INVENTION: Thioredoxin and Thioredoxin-Like Molecules
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 926-6200
TELEFAX: (415) 854-3713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 10-FEB-199
CLASSIFICATION: 435
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MEDIUM TYPE: 3.5" diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Weil, Gotshal & Manges
STREET: 2882 Sand Hill Road, Ste.
CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U
ZIP: 94025
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  ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 47; DB 1; Length 106; ilarity 100.0%; Pred. No. 4.5; Conservative 0; Mismatches 0; Indels
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VENTION: An oxido reductase enzyme system

VENTION: obtainable from P. chrysogenum, the set of genes

VENTION: encoding the same and the use of oxido reductase enzyme

VENTION: systems or genes encoding the same for increasing antibiotic

VENTION: production
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Kattevilder, Alfred
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Genetics Institute, Inc.
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          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/921,848
FILING DATE: 19920728
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/652,531
FILING DATE: 06-FEB-1991
PRIOR APPLICATION NUMBER: US 07/745,382
APPLICATION NUMBER: US 07/745,382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTMARE: PACENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/745,382
FILING DATE: 1991.0814
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/652,531
FILING DATE: 06-FEB-1991
ATTORNEY/AGENT INFORMATION:
NEMBER: CEST. LINEAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 5292646
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 22:
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CITY: Ca.
COUNTRY:
                                                                                                                                                                                                                                                                                                   STREET: 0, STREET: Cambridge CITY: Cambridge CTATE: Massachusetts
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REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI5188A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
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LENGTH: 109 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: McCoy, John
APPLICANT: LaVallie, Edward
TITLE OF INVENTION: Peptide and Protein Fusions To
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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ZIP: 02140
FILING DATE:
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o. 5292646
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                                                                                                                                                                                                                                                  Floppy disk
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14-AUG-1991
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Pred. No. 4.6;
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ATTORNEY/AGENT INFORMATION:

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                 Query Match
Best Local Similarity
Watches 6; Conserve
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US-08-165-301A-22
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                                                                                                       US-08-165-301A-22
                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI 5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
Matches 6; Conserv
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/165,301A
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
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TELEPAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Grant, Kathleen
APPLICANT: Lavallie, Edward R.
APPLICANT: LAVALLIE, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES,
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: protein
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                                                                                                                    LENGTH: 109 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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APPLICANT: DiBlasio-Smith, Elizabeth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: M
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TYPE: AMINO ACID
TOPOLOGY: linear
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   1 WCGPCK 6
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T: U.S.A.
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87 CambridgePark Drive
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                                                  100.0%;
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Pred. No.
                                                  Score 47; DB
Pred. No. 4.6;
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                                                                   DB 1; Length 109;
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US-08-165-301A-28
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Patent No.
                                                                                                                                                                                                            Sequence 28, Appli
Patent No. 5646016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/165,301A
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 33,544
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI 5188D
TELEPAN: (617) 876-1170
TELEPAN: (617) 876-5170
                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local Similarity
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                           APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: THIOREDOXIN,
TITLE OF INVENTION: THIOREDOXIN,
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (617) 876-585
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS LENGTH: 109 amino aci
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                          APPLICANT: McCoy, John APPLICANT: DiBlasio-Smith,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: McCoy, John APPLICANT: DiBlasio-Sm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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STREET:
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                  ADDRESSEE:
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5. 5646016
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87 CambridgePark Drive
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87 CambridgePark Drive
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                Genetics Institute,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 47; DB 100.0%; Pred. No. 4.6;
                                                                    THIOREDOXIN, THIOREDOXIN-L
THIOREDOXIN-LIKE MOLECULES
                                                                                   PEPTIDE AND PROTEIN FUSIONS TO THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Edward R.
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COUNTRY:

Massachusetts: U.S.A.

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      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/810,436
FILING DATE: 04-MAR-1997
CLASSIFICATION #35
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/165,301
APPLICATION NUMBER: 08/165,301
FILING DATE: 10-DEC-1993
ATTORNEY/AGENT IMFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REGISTRATION NUMBER: 33,544
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/165,301A
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
THE TRANSPORTATION NUMBER: 33,544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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(INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
APPLICANT: LAVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
NUMBER OF SEQUENCES: 29
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                     CITY: Cambridge
STATE: Massing
                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A. ZIP: 02140
REFERENCE/DOCKET NUMBER:
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XGY: linear
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GI 5188D
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RESULT 17
US-08-810-436-26
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                                                                  US-08-810-436-26
                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/810,436
FILING DATE: 04-MAR-1997
CLASSIFICATION 04-MAR-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/165,301
FILING DATE: 10-DEC-1993
ATTORNEY/AGENT: INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
PRICE TRANSION NUMBER: 33,544
PRICE TRANSION NUMBER: 33,544
Best Local Similarity
Matches 6; Conserv
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                              Query Match
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith,
APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward
                                                                                                                              IELEFAX: (617) 876-585
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                              MOLECULE TYPE: protein
                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                  TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 02140
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Local Similarity 100.0%;
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87 CambridgePark Drive
                                                                                                                                109 amino acids
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LaVallie, Edwar
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876-5851
NO: 26:
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TO NO: 22:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , Edward R.
PEPTIDE AND PROTEIN FUSIONS TO
THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
                100.0%; Score 47; DB 100.0%; Pred. No. 4.6;
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 Mismatches
                                DB 2;
                                Length 109;
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RESULT 18 US-08-810-436-28

Patent No.

8, Application US/08810436 6143524 밁

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RESULT 19
US-09-485-885-17
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                                                                                        Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Celisse, Anne-Marie Eva Fernande
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
                                                                                                                                                                                                                                                                                                                                                                         Sequence 17, Application US/09485885 Patent No. 6342224
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Best Local Similarity
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/810,436
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APPLICANT: McCoy,
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
                                                                         FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (617) 876-1
TELEPAX: (617) 876-585
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: ITTILE OF INVENTION: TITLE OF INVENTION: TOTAL TOTAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/165,301 FILING DATE: 10-DEC-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith, Elizabeth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 04-MA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
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87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             : (617) 876-5851
(617) TO NO: 28:
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LaVallie, Edward R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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SEQ ID NO 19911

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                                                                                                                                                                                                                                                                       RESULT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: amino acid of thioredoxin US-09-166-966E-10
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LENGTH: 109
TYPE: PRT
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APPLICANT: CHEN, DAVID CHANHAN
APPLICANT: HU, WIEN-TAI
APPLICANT: CHEN, YUN-JU
APPLICANT: HSEU, TZONG-HSIUNG
APPLICANT: HSEU, TZONG-HSIUNG
                                                                                  GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
CURRENT FILING DATE: 1999-02-12
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SEQ ID NO 10
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               PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: TW 86114750 PRIOR FILING DATE: 1997-10-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: CHIMERIC PROTEINS WITH A CELLULOSE BINDING FILE REFERENCE: 32350-150960 CURRENT APPLICATION NUMBER: US/09/166,966E CURRENT FILING DATE: 1998-10-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: PCT, PRIOR FILING DATE: 1998-08-17 PRIOR APPLICATION NUMBER: GB: PRIOR FILING DATE: 1997-08-22
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NUMBER OF SEQ ID NOS: 28208
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NAME/KEY: thioredoxin
LOCATION: 1..109
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TYPE: PRT
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Pred. No. 4.6
0; Mismatches
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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RESULT 23
PCT-US94-14179-26
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Best Local Similarity
Watches 6; Conserve
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US-09-248-796A-19911
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PCT-US94-14179-22
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                                                Sequence 26, Application PC/TUS9414179 GENERAL INFORMATION:
                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                     TELEPAX: (617) 876-5851
TELEFAX: (617) 876-5851
TELEFAX: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: DiBlasio-Smith, Elizabeth
APPLICANT: Grant, Kathleen
APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith, Elizabeth
APPLICANT: Grant, Kathleen
                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Melhert, M. C.
REGISTRATION NUMBER: 33,544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/14179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                           TYPE: amino acid
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Pred. No. 4.6;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                         5188D
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RESULT 24
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CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GIT
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acide
TYPE: amino acide
TYPE: amino acide
TYPE: amino acide
TYPE: amino acide
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GENERAL INFORMATION:
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TITLE OF INVENTION: THI
TITLE OF INVENTION: THI
TITLE OF INVENTION: THI
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/14179
                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: McCoy, John APPLICANT: DiBlasio-Smith,
                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                        CITY: Cambridge
STATE: Massachus
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                    APPLICATION NUMBER: FILING DATE:
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                                                                                                                                                                                                                 COUNTRY:
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CLASSIFICATION:
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87 CambridgePark Drive
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IVENTION: PEPTIDE AND PROTEIN FUSIONS TO

IVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES,

IVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                   U.S.A.
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                                         PCT/US94/14179
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US-09-134-001C-3859
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US-09-264-419C-6
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          Sequence 3859, Application US/09134001C

Patent No. 6380370

GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: REIDERWIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC-007
CURRENT APPLICATION NUMBER: US/09/134,001C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 3859
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Best Local S
Matches 6
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CURRENT FILING DATE: 1999-03-08
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 6
LENGTH: 111
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: No. 6174682el Thioredoxin Family Active Site Molecules and Uses TITLE OF INVENTION: Therefor FILE REFERENCE: MNI-076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 28:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 109 amino acids
TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                             WCGPCK 38
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Query Match
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S
                                                                                                ; ORGANISM: Proteus mirabilis US-09-543-681A-4397
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US-09-882-835-5
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                                                                                                                               NUMBER OF SEQ ID NOS: 8344
SEQ ID NO 4397
LENGTH: 122
TYPE: PRT
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APPLICANT: GARY BRETON
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Best Local Similarity
Matches 6; Conser
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Patent No. 6462187
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                                   Matches
                                                  Query Match
Best Local :
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CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/211,673
PRIOR FILING DATE: 2000-06-15
NUMBER OF SEQ ID NOS: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Bandaru, Rajasekhar
TITLE OF INVENTION: 22109, A NOVEL HUMAN THIOREDOXIN FAMILY
TITLE OF INVENTION: MEMBER AND USES THEREOF
FILE REFERENCE: 10448-063001
                                                                                                                                                                                             TITLE OF INVENTION: NUCLEIC ACID AND AMINO TITLE OF INVENTION: DIAGNOSTICS AND THERAI FILE REFERENCE: 2799.1002-001 CURRENT APPLICATION NUMBER: US/09/543,681A CURRENT FILING DATE: 2000-04-05 PRIOR APPLICATION NUMBER: US 60/128,706 PRIOR EILING DATE: 1999-04-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
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   1 WCGPCK 6
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                              100.0%; ilarity 100.0%; Conservative 0;
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100.0%; Pred. No.
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                                 Score 47; DB
Pred. No. 5;
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                                                                  DB 2;
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RESULT 31
US-09-272-342B-5
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Best Local Similarity
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PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 8017
                                                                                                  Sequence 5, Application US/09272342B Patent No. 6294341
                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 9663
LENGTH: 125
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                                                                                   GENERAL INFORMATION:
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APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local Similarity
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Patent No. 6812339
                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Human
-09-949-016-9663
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CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILLING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILLING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILLING DATE: 2000-09-08
NUMBER: 622-72, WG 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
APPLICANT: KIM, SUNG-HOU
APPLICANT: RYU, JAE-RYEON
TITLE OF INVENTION: METHOD FOR DETECTING A SUBSTANCE HAVING AN ACTIVITY TO
                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: GATY Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
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                                                                                                                                                                                                                            WCGPCK 56
                                                           YU, YEON-GYU
                                                                                                                                                                                                                                                                                                         100.0%; Score 47; DB 2; Length 125; ilarity 100.0%; Pred. No. 5.1; Conservative 0; Mismatches 0; Indels
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Pred. No. 5.
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SOPTWARE: PatentIn Ver.
SEQ ID NO 5
LENGTH: 134
TYPE: PRT
ORGANISM: Homo sapiens
US-09-272-342B-5
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US-08-961-083-40
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                                                                                          US-08-961-083-40
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GENERAL INFORMATION:
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                               Matches
                                                            Query Match
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                                                                                                                                                     TELEFAX: (301) 303-00-101 INFORMATION FOR SEQ ID NO: 40: SEQUENCE CHARACTERISTICS: 1-ENGTH: 137 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/272,342B
CURRENT FILING DATE: 1999-03-19
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Brookes, A. Anders
REGISTRATION NUMBER: 36,373
REFERENCE/DOCKET NUMBER: PB34
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                         MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM:
                                                                                                                         TOPOLOGY:
                                                                                                                                 STRANDEDNESS:
                                                                                                                                                    TYPE: amino acid
                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 20850
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                                              Local
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                              Similarity 6; Conser
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   WCGPCK 6
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ilarity 100.0%;
Conservative 0
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                              100.0%; Score 47; DB larity 100.0%; Pred. No. 5.5; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Streptococcus pneumoniae Antigens and Vaccines 452
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.50 inch, 1.4Mb storage
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                                                          DB 2; Length 137;
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RESULT 34
US-09-765-271-40
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                                                                                                                                                                                                                                            Sequence 40, Application US/09765271
Patent No. 6887663
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 40, Application US/09536784
Patent No. 6573082
Patent NIFORMATION:
APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
NUMBER OF SEQUENCES: 452
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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INFORMATION FOR SEQ ID NO: 40:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/961,083
FILING DATE: OCT-30-1997
ATTORNEY/AGENT INFORMATION:
NAME: MICHELLE S. MARKE
REGISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB340P3
TELECOMMUNICATION INFORMATION:
TELECHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity hes 6; Conserv
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APPLICATION NUMBER: US/09/536,784
FILING DATE: 30-Oct-1997
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
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MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 40:
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                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 452
                                                                                                                                                                                                                                                                                                                                                                                                                                    61
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                                                                                  ZIP:
                                                                                                                        CITY: Rockville
STATE: Maryland
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 137 amino acids
TYPE: amino acid
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                                                                                                       COUNTRY:
                                                                                                                                                                                   ADDRESSEE: Human Genome Sciences, Inc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 40, Application US/09:
Patent No. 692930
GENERAL INFORMATION:
APPLICANT: Choi et. al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
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SEQUENCE CHARACTERISTICS:
LENGTH: 137 amino acids
TYPE: amino acid
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 09/536,784
FILING DATE: «Unknown»
APPLICATION NUMBER: 08/961,083
FILING DATE: OCT-30-1997
ATTORNEY/AGENT INFORMATION:
NAME: Michelle S. Marks
REGISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB340P3
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8512
TELEPHONE: (301) 309-8512
                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/961,083
APPLICATION NUMBER: 09/961,083
FILING DATE: 0CT-30-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lin J. Hymel
REGISTRATION NUMBER: 45,414
REFERENCE/DOCKET NUMBER: PB340P2C2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage COMPUTER: Dell Latitude C610

OPERATING SYSTEM: Windows 2000

SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/765,272A

FILING DATE: 22-Jan-2001

CLASSIFICATION: Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 45 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 WCGPCK 66
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                                    TELEPHONE: (301)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Human Genome Sciences, Inc.
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100.0%; Pred. No. 5.5;
7ative 0; Mismatches
                                                              (301)
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                                      309-8439
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US-09-328-352-5360
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                                                                   ; LOCATION: 1..109
; OTHER INFORMATION: Combined amino acid of thioredoxin and an artificial sequence
US-09-166-966E-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          밁
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// ORGANISM: Acinetobacter baumannii
US-09-328-352-5360

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Matches
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SEQ ID NO 5360
LENGTH: 145
TYPE: PRT
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APPLICANT: CHEN, DAVID CHANHAN
APPLICANT: HU, NIEN-TAI
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APPLICANT: GATY L. Breton et al.
APPLICANT: TITLE OF INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 5360, Application US/09328352 Patent No. 6562958
Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                SEQ ID NO 8
LENGTH: 159
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Patent No. 6407208
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Best Local Similarity
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                                                                                                                                                                                                              APPLICANT: CHEN, YUN-JU
APPLICANT: CHEN, TYN-JUNG
APPLICANT: HSEU, TZONG-HSIUNG
TITLE OF INVENTION: CHIMERIC PROTEINS WITH A CELLULOSE BINDING DOMAIN
FILE REFERENCE: 32350-150960
CURRENT APPLICATION NUMBER: US/09/166,966E
CURRENT FILING DATE: 1998-10-06
PRIOR APPLICATION NUMBER: TW 86114750
PRIOR PILING DATE: 1997-10-08
NUMBER OF SEQ ID NOS: 11
                                                                                                               PEATURE:
NAME/KEY: thioredoxin
                                                                                                                                                   ORGANISM: Unknown
                                                                                                                                                                      TYPE: PRT
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     Conservative
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   Score 47; DB 2; Length 159; Pred. No. 6.3; Indels
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RESULT 39
US-08-775-978-3
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US-08-775-978-1
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                                                                                                                                                            Sequence 3, Application US/08775978
Patent No. 5831049
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                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                          GENERAL INFORMATION:
APPLICANT: HILLMAN, JENNIFER L.
APPLICANT: GOLI, SURYA K.
TITLE OF INVENTION: NOVEL HUMAN THIOREDOXIN
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: HILLMAN, JENNIFER L.
APPLICANT: GOLI, SURYA K.
TITLE OF INVENTION: NOVEL HUMAN THIOREDOXIN
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/775,978
FILING DATE: TO BE ASSIGNED
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                     IMMEDIATE SOURCE:
CLONE: 1430906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 3174 Por
CITY: Palo Alto
STATE:
            STREET: 3174 POI
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single
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                          ADDRESSEE: Incyte Pharmaceuticals, Inc
STREET: 3174 Porter Drive
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COUNTRY:

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US-09-741-243C-4
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                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/741,243C
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: 60/171,785
PRIOR FILING DATE: 1999-12-22
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FeatSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 167
                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Crae, John
APPLICANT: Rice, John
APPLICANT: Sevala, Veeresh
APPLICANT: Stewart, Sandy
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4, Application US/09741243C Patent No. 6399352
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                                                           Best Local Similarity
Matches 6; Conserv
                                                                                          Query Match
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APPLICANT: Stewart, Sandy
TITLE OF INVENTION: NOVEL PLANT PORPHOBILINGEN AND FUSION
TITLE OF INVENTION: PROTEIN THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 166 amino acids
                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 2022US
                                                                                                                                                       ORGANISM: Unknown FEATURE:
                                                                                                                                                                                      TYPE: PRT
                                                                                                                                         OTHER INFORMATION: Thioredoxin functional fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/775,978 FILING DATE: TO Be Assigned PRIOR APPLICATION DATA:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
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nes 6; Conserv
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32
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                                1 WCGPCK 6
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WCGPCK 37
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                                                                           Score 47; DB 2; Length 167; Pred. No. 6.5;
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RESULT 42
US-09-270-767-47690
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US-09-270-767-32473
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Patent No. 6703491
GRMERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7336-094
CURRENT APPLICATION NUMBER: US/09/270,767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins elle Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 47590
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Matches
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NUMBER OF SEQ ID NOS: 62517
SOFTWARE: PATENTIN VEr. 2.0
SEQ ID NO 32473
LENGTH: 170
                                                                                                                                                                                                                                                                                                                                             Best
                                                                                 GENERAL INFORMATION:
APPLICANT: YU, YEON-GYU
APPLICANT: KIM, SUNG-HOU
APPLICANT: RYU, JAE-RYEON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                      Sequence 6, Application US/09272342B Patent No. 6294341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 47690, Application US/09270767 Patent No. 6703491
                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
            TITLE OF INVENTION: METHOD FOR DETECTING A SUBSTANCE HAVING AN ACTIVITY TITLE OF INVENTION: INHIBIT HIV INFECTION USING IMMUNOASSAY AND VARIANT TITLE OF INVENTION: PROTEIN USED FOR SAID METHOD FILE REFERENCE: 2901-0125-0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT ORGANISM: Drosophila melanogaster
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                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Xaa means any amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
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 APPLICATION
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NUMBER: US/09/272,342B
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100.0%; Pred. No. 6.
tive 0; Mismatches
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Pred. No.
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Best Local Similarity
Thes 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/583,110
CURRENT FILING DATE: 200-05-26
PRIOR APPLICATION NUMBER: US 09/107,433
PRIOR FILING DATE: 1998-06-30
PRIOR APPLICATION NUMBER: US 60/085,131
PRIOR FILING DATE: 1998-05-12
PRIOR APPLICATION NUMBER: US 60/051,553
PRIOR FILING DATE: 1997-07-02
NUMBER: OF SEQ ID NOS: 5322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 5222
LENGTH: 185
                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lynn Doucette-Stamm et al.
TITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Relating to Streptococcus
TITLE OF INVENTION: Pneumoniae for Diagnostics and Therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT FILING DATE: 1999-03-19
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: homo sapiens and HIV hybrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 177
                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                              COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5222, Application US/09583110
5. 6699703
                                                                                                                                                                                                                                                                                                                                                            80 WCGPCK 85
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COMPUTER: <Unknown>
OPERATING SYSTEM: <Unknown>
                                                                                          STATE: Massachusetts
                                                                                                          STREET: 100 Beaver Street CITY: Waltham
                                                                             COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 47; DB 2; Length 177; ilarity 100.0%; Pred. No. 6.8; Conservative 0; Mismatches 0; Indels
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Pred. No. 7
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Query Match

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                                                                                                                                                                                                                                                                                        APPLICANT: CHEN, YUN-JU
APPLICANT: HSEU, TZONG-HSIUNG
APPLICANT: HSEU, TZONG-HSIUNG
TITLE OF INVENTION: CHIMERIC PROTBINS WITH A CELLULOSE BINDING DOMAIN
FILE REFERENCE: 32350-150960
CURRENT FILING DATE: 1998-10-06
CURRENT FILING DATE: 1998-10-06
PRIOR APPLICATION NUMBER: TW 86114750
PRIOR FILING DATE: 1997-10-08
NUMBER OF SEQ ID NOS: 11
SEQ ID NO 11
SEQ ID NO 11
SEQ ID NO 11
SEQ ID NOS: 209
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Matches
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GENERAL INFORMATION:
                                                                                       FEATURE:
NAME/KEY: thioredoxin
LOCATION: 1..109
NAME/KEY: Cellose-binding domain
LOCATION: 162..197
NAME/KEY: RGD
LOCATION: 204..206
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: CHEN, DAVID C
  OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (781)893-8277
INFORMATION FOR SEQ ID NO: 5104:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Unknown
                                            OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/107,433
FILING DATE: 30-Jun-1998
FRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 60/ 085131
FILING DATE: May 12, 1998
APPLICATION NUMBER: 60/051553
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40, 489
REFERENCE/DOCKET NUMBER: GTC-011
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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LOCATION: (B) LOCATION 1...194
SEQUENCE DESCRIPTION: SEQ ID NO: 5104:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: protein HYPOTHETICAL: YES ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 194 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (781)893-5007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 47; DB: llarity 100.0%; Pred. No. 7.4; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DAVID CHANHAN
Combined amino acid sequence of thioredoxin, artificial sequence, cellulose-binding domain, and artificial sequence; the two artificial sequences that two artificial sequences flanking the cellulose-binding domain contain restriction sites
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PCT-US92-05707-6
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Matches 6; Conserv
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TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 210 amino acids
       TITLE OF INVENTION: Hematopoietic Growth Factor Derived from TITLE OF INVENTION: Tlymphocytes and Methods of Use Therefor NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
TELEPAX: (312) 474-0448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 906866
FILING DATE: 01 July 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
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les 6; Conserv
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REGISTRATION NUMBER: 30,060
REFERENCE/DOCKET NUMBER: 276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    106 WCGPCK 111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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Marshall, O'Toole, Gerstein, Murray & Bicknell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-JAN-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US/08/180,371
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); Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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Query Match
Best Local Similarity
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US-08-778-717-17
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMBER: PCT/US92/05707
FILING DATE: 19920707
CILASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Gruber, Lewis S.
REGISTRATION NUMBER: 30,060
REFERENCE/DOCKET NUMBER: 27620/30933
             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: BC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/778,717
FILING DATE: 12-DEC-1996
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 352225/1995
FILING DATE: 28-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17, Application US/08778717 Patent No. 6602689
                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:

APPLICANT: UENO, EIICHI

APPLICANT: WOBUYUKI, FUJII

APPLICANT: OKADA, MASAHISA

TITLE OF INVENTION: FUSED DAA SEQUENCE, FUSED PROTEIN

TITLE OF INVENTION: EXPRESSED FROM SAID FUSED DNA SEQUENCE AND METHOD FOR

TITLE OF INVENTION: EXPRESSING SAID DNA SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 346-5750
TELEPAX: (312) 984-9740
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: AMINO ACID
TOPOLOGY: line
OLECTION
                                                                                                                                                                                                     ZIP: 22202
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: I
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STREET:
CITY: C
                                                                                                                                                                                                                                                                                       ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, PALLA L. ..... ADDRESSEE: P.C. STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400 CITY: ARLINGTON
                                                                                                                                                                                                                                                 COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            106 WCGPCK 111
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: Illinois
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Two First National Plaza,
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                                                                                                                                                                                                                                                                                                                                                             OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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NAME: OBLON, NORMAN F.

REGISTRATION NUMBER:

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                                                                       SOFTWARE: Patentin Release #1.0, Ver CURRENT APPLICATION DATA: APPLICATION UMBER: US/07/745,382 FILING DATE: 19910814 CLASSIFICATION: 435 PRIOR APPLICATION UMBER: US 07/652,531 PRIOR DATE: 06-FEB-1991 ATTORNEY/AGENT INFORMATION: NAME: CGETT, LUAND REGISTRATION NUMBER: 31,822 REFERENCE/DOCKET NUMBER: 31,822 REFERENCE/DOCKET NUMBER: G15188A TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 876-1170 TELEPAX: (617) 876-5851 INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 14, Application US/07745382
Patent No. 52770181
GENERAL INFORMATION:
APPLICANT: McCoy, John
APPLICANT: LaVallie, Edward
APPLICANT: LaVallie, Edward
TITLE OF INVENTION: Peptide and Protein Fusions To
TITLE OF INVENTION: Thioredoxin and Thioredoxin-Like Molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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SEQUENCE CHARACTERISTICS:
LENGTH: 265 amino acids
               SEQUENCE CHARACTERISTICS:
LENGTH: 296 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: RECOMBINANT
PUBLICATION INFORMATION:
AUTHORS: NOBUYUKI FUJII ET AL,
AUTHORS: NOBUYUKI FUJII ET AL,
TITLE: FUSED DNA SEQUENCE, FUSED PROTEIN EXPRESSED
TITLE: FROM SALD FUSED DNA SEQUENCE AND METHOD OF
TITLE: EXPRESSING SAID FUSED PROTEIN
RELEVANT RESIDUES IN SEQ ID NO: 17: FROM 1 TO 265
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
TELEPHAX: 703-413-2220
                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: |
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: Mass
COUNTRY: U.
ZIP: 02140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 265 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          50 WCGPCK 55
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                                                                                                                                                                                                                                                                                                                                                                                                   Release #1.0, Version #1.25
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RESULT 52
US-08-115-680-4
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                                                                                                                                                                                                                       ; TOPOLOGY: linear; MOLECULE TYPE: protein US-07-921-848-14
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US-07-921-848-14
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Best Local Similarity
watches 6; Conserva
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                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC COMPACTIBLE
COMPUTER: IBM PC COMPACTIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PACENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/921,848
FILING DATE: 19920728
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/652,531
FILING DATE: 06-PEB-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/745,382
FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: CGETY, LUADIN
REGISTRATION NUMBER: G15188A
TELECOMMUNICATION INFORMATION:
TELESPHONE: (617) 876-1110
TETESPENCE, JOCKET NUMBER: G15188A
TELECOMMUNICATION INFORMATION:
TELESPHONE: (617) 876-1110
                                                                                                                                               Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 5292646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14, Application US/07921848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 2: CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Peptide and Protein Fusions To
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: McCoy, John APPLICANT: LaVallie, E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 87 Cambridge
CITY: Cambridge
STATE: Massachusetts
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                           32 WCGPCK 37
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                                                                                                                                                                                                                                                                          4: 296 amino acids
AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                    : (617) 876-5851
(617) 876-5851
TD NO: 14:
                                                                                                                                               100.0%; Score 47; DB ilarity 100.0%; Pred. No. 10; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                    DB 1; Length 296;
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Sequence 4, Application US/08115680 Patent No. 5437863

GENERAL INFORMATION:

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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/941,372
APPLICATION NUMBER: US 07/941,372
FILING DATE: 02-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: INDIAUSA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 540-9206
TELEPAX: (215) 540-9206
TELEPAX: (215) 540-9218
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
INVERTED: 366 Entic 36146
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4, Application US/07941372 Patent No. 5460810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Williams, David A.
APPLICANT: Clark, Steven C.
ZIP: 194/
COMPUTER READABLE FORM:
COMBUTUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION NUMBER: US/07/941,372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Williams, David A.
APPLICANT: Clark, Steven C.
TITLE OF INVENTION: Method of Treating Cell Damage or
TITLE OF INVENTION: Depletion
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Method of TITLE OF INVENTION: Depletion NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/115,680
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Howson and Howson
STREET: Spring House Corporate Cntr, P.O. Box 457
CITY: Spring House
                                                                                                                                                                                                         ADDRESSEE: Howson and Howson
STREET: Spring House Corporate Center,
CITY: Spring House
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 296 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity tes 6; Conserv
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US-08-165-301A-14
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                                                                          US-08-165-301A-14
                                                                                                                                                                                      APPLICATION NUMBER: US/08/165
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI 5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 564601
Query Match 100.0%; Score 47; DB Best Local Similarity 100.0%; Pred. No. 10; Matches 6; Conservative 0; Mismatches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (215) 540-5818 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 296 amino acids
TYPE: AMINO ACID
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REGISTRATION INVESE: 31,215
REFERENCE/DOCKET NUMBER: INI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 540-9206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION: NAME: Bak, Mary E.
                                                                                                                                                  SEQUENCE CHARACTERISTICS LENGTH: 296 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: LAVAllie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
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CORRESPONDENCE ADDRESS:
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                                                                                          TOPOLOGY: linear MOLECULE TYPE: protein
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GY: linear
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RESULT 55
US-08-810-436-14
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PCT-US93-08247-4
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                                                                                                                                                  Sequence 4, Application PC/TUS9308247 GENERAL INFORMATION:
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/165,301
PILING DATE: 10-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: Melhert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI 5188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/810,436
FILING DATE: 04-MAR-1997
                                                APPLICANT: Genetics Institute, Inc., , APPLICANT: 87 CambridgePark Drive, , APPLICANT: Cambridge, MA 02140, USA TITLE OF INVENTION: Method of Treating TITLE OF INVENTION: Depletion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
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APPLICANT: DiBlasio-Gmith, Eli:
APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                  NUMBER OF SEQUENCES:
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COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
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  ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 47; DE ilarity 100.0%; Pred. No. 10; Conservative 0; Mismatches
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Howson and Howson
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04-MAR-1997
                                                                         Method of Treating Cell Damage or
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          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/14179
FILING DATE:
CLASSIFICATION:
NAME: Meinert, M. C.
NAME: Meinert, M. C.
REGISTRATION UMBER: 33,544
REGISTRATION UMBER: 33,544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 14, Application PC/TUS9414179 GENERAL INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION UMMBER: US 07/941
APPLICATION UMMBER: US 07/941
FILING DATE: 02-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: IND1.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 540-9206
TELEPHONE: (215) 540-5818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (215) 540-581
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith,
APPLICANT: Grant, Kathleen
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08247
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                CITY: Cambridge
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid TOPOLOGY: linear
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REFERENCE/DOCKET NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        296 amino acids
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LaVallie, Edwar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Edward R.
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GI 5188D
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TELECOMMUNICATION INFORMATION:

TELEPHONE:

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TELEFAX: (617) 876-5851;
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 296 amino acids
TYPE: amino acid
TYPE: mino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
PCT-US94-14179-14
                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 1011
SOPTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1011
LENGTH: 355
TYPE: PRT
ORGANISM: Homo sapiens
US-10-012-896-1011
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RESULT 59
US-08-683-007A-2
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Best Local Similarity
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US-10-012-896-1011
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                                                                                                                                                                                                                                                                                                  APPLICANT: Meagher, Madeleine Joy
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427027
CURRENT APPLICATION NUMBER: US/10/012,896
CURRENT FILING DATE: 2001-12-10
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APPLICANT:
APPLICANT:
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APPLICANT: Dillon, Davin C.
Tennife
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                                                                                           1 WCGPCK 6
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Carter, Darrick
Li, Samuel X.
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Fanger, Gary R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Henderson, Robert A.
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Hepler, William T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Retter, Marc W. Stolk, John A.
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Kalos, Michael D.
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                                                                                                                            Conservative
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SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 2
LENGTH: 467
TYPE: PRT
ORGANISM: Mus musculus
US-09-513-442-2
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TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

US-08-683-007A-2
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                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Ikemoto, mitsushi
TITLE OF INVENTION: SPARC Fusion Protein and
TITLE OF INVENTION: Same
FILE REFERENCE: HIRAKI-04218
                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/09513442 Patent No. 6387664
                              Query Match 100.0%; Score 47; DI
Best Local Similarity 100.0%; Pred. No. 15;
Matches 6; Conservative 0; Mismatches
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Best Local S
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APPLICANT: No. 5858724y, Robert E
APPLICANT: Domanico, Michael
APPLICANT: Yaeger, Keith
APPLICANT: Yaeger, Keith
APPLICANT: Kroeker, Warren
TITLE OF INVENTION: Recombinant Rabbit Tissue Factor
NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/513,442
CURRENT FILING DATE: 2000-02-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 608-251-9166
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27386
REFERENCE/DOCKET NUMBER: 74
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-251-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 16 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/683,007A
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100.0%; Pred. No. 14;
tive 0; Mismatches
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                                                     DB 2; Length 467;
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RESULT 61 US-08-260-582-77

Application US/08260582

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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: McCoy, John M.
APPLICANT: Lu, Zhijian
TITLE OF INVENTION: METHOD OF DETECTING LIGAND
TITLE OF INVENTION: INTERACTIONS
                                                                                                                                                                                                                                                                                                                          Sequence 77, Application PC/TUS9505471 GENERAL INFORMATION:
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Best Local Similarity
Matches 6; Conserv
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INFORMATION FOR SEQ ID NO: 77:
SEQUENCE CHARACTERISTICS:
LENGTH: 500 amino acids
                                                                                     APPLICATION NUMBER: PINFORMATION FOR SEQ ID NO:
                                                                                                                          TITLE OF INVENTION: METHOD OF DETECTING LIGAND INTERACTIONS
NUMBER OF SEQUENCES: 76
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/00
FILING DATE: 16-JUN-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
COMPUTER: FORDS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CORRENT APPLICATION DATA:
CORRENT APPLICATION DATA:
CORRENT APPLICATION UNDERS. 10/08/06/06
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CORRESPONDENCE ADDRESS:
                                                               SEQUENCE CHARACTERISTICS:
                        LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Meinert, M. C.
REGISTRATION NUMBER: 31,544
REFERENCE/DOCKET NUMBER: GI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
LENGTH: 500 amino acids
TYPE: amino acid
TOPOLOGY: linear
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87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             U.S.
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                                                                                                            PCT/US95/05471
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RESULT 64
US-09-741-243C-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 63
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-057-531A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Lyon, Jeffrey A.
APPLICANT: Lyon, Jeffrey A.
APPLICANT: Lyon, Jeffrey A.
APPLICANT: Angov, Evelina
TITLE OF INVENTION: Isolation and Purification of P. falciparum Merozoite
TITLE OF INVENTION: Protein-142 Vaccine
FILE REFERENCE: 003/241/SAP
CURRENT APPLICATION NUMBER: US/10/057,531A
CURRENT FILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: US 60/264,535
PRIOR APPLICATION NUMBER: US 60/347,564
PRIOR FILING DATE: 2001-10-26
PRIOR FILING DATE: 2001-10-26
                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Crawford Jr., John Milton
APPLICANT: Rice, John
APPLICANT: Sevala, Veeresh
APPLICANT: Sevala, Veeresh
APPLICANT: Stewart, Sandy
TITLE OF INVENTION: NOVEL PLANT PORPHOBILINOGEN AND FUSION
TITLE OF INVENTION: PROTEIN THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                  SOFTWARE: FastSEQ for Windows Version SEQ ID NO 2
LENGTH: 551
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Apple Macintosh Microsoft Word 6.0 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/10057531A Patent No. 6855322
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/09741243C
                                                                                                                                                                        FILE REFERENCE: 2022US
CURRENT APPLICATION NUMBER: US/09/741,243C
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: 60/171,785
PRIOR FILING DATE: 1999-12-22
                                                                                                                                                        NUMBER OF SEQ ID NOS: 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Artificial sequence
OTHER INFORMATION: A plant thioredoxin-porphobilinogen
                      ORGANISM: Artificial Sequence FEATURE:
                                                                    TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
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Local Similarity 100.0%;
hes 6; Conservative 0
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ilarity 100.0%; Pred. No. 16
Conservative 0; Mismatches
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Pred. No.
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                                     US-09-626-589-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-626-589-3
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                                                                                                                                                                SEQ ID NO 3
LENGTH: 824
TYPE: PRT
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Best Local S
Matches 6
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APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: AUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/626,589
CURRENT FILING DATE: 2000-07-27
NUMBER OF SEQ ID NOS: 6
SOFTWARE: FASTSEO for ""."
EG TO ""."
                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application US/09626589 Patent No. 6326164
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Best Local Similarity
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LENGTH: 580
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Query Match
                                                                                                                                                                                                                                                                                 APPLICANT: Rice, John
APPLICANT: Kloti, Andreas
APPLICANT: Crawford, John
APPLICANT: Lanning, Beth
APPLICANT: Lanning, Beth
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE
TITLE OF INVENTION: INDENTIFICATION OF MODULATORS OF
TITLE OF INVENTION: SYNTHASE ACTIVITY
FILE REFERENCE: 2037 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: UPRIOR FILING DATE: 1998-07-
NUMBER OF SEQ ID NOS: 33142
                                               FEATURE:
OTHER INFORMATION: Residues 1-165 are from the thioredoxin sequence
OTHER INFORMATION: found in the vector pET32 supplied by No. 6326164ac
OTHER INFORMATION: Residues 166-824 represent the tDXPS sequence from
OTHER INFORMATION: Arabidopsis shown in SEQ ID NO:2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Pseudomonas aeruginosa
                                                                                                                                      ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    504 WCGPCK 509
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                                                                                                                                                                                                                   FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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1998-02-18
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100.0%;
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Pred. No.
Score 47;
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DB 2;
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Length 824;
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Query Match
Best Local Similarity
"""hes 6; Conserv
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US-10-237-551-92
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APPLICANT: Hosken, Nancy A.
APPLICANT: Hosken, Nancy A.
APPLICANT: Parsons, Joseph M.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AN
TITLE OF INVENTION: TREATMENT OF HERPES SIMPLEX VIRUS INFECTION
FILE REFERENCE: 210121.538C3
CURRENT APPLICATION NUMBER: US/10/237,551
CURRENT FILING DATE: 2002-09-06
NUMBER OF SEQ ID NOS: 254
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 92
LENGTH: 858
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 600,
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Matches 6; Conserv
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              APPLICATION NUMBER: US/08/
FILING DATE: 19-NOV-1997
CLASSLFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 01-OCT-1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                       COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                           ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                APPLICANT: Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Cech, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                              STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                COUNTRY:
 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
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682151
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Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/08974549A
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100.0%; Pred. No.
                                                                                                                      US/08/974,549A
US 08/844,419
                                                   08/724,643
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                                                                                                                                                         Version #1.30
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FILING DATE: 18-APR-1997

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RESULT 69
US-08-912-951-314
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                                                                                                                                                                                                                                                            Sequence 314, Application US/08912951 Patent No. 6475789
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                           Patent No. 6475789
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                          APPLICANT:
                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph Teo
REGISTRATION NUMBER: 36,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/17885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATE: 06-PAR.
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/854,050
APPLICATION NUMBER: 09-WAX-1997
PATE: 09-WAX-1997
                                                                          TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
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APPLICATION NUMBER:
                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein
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OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
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                                             ADDRESSEE:
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IT: Ling.

NT: Nakamura, Tol.

NT: Chapman, Karen B.

ANT: Morin, Gregg B.

ANT: Marley, Calvin

CANT: Andrews, William H.

OF INVENTION: HUMAN TELOMERASE CATALYTIC SUBUNIT: DIAGNOSTIC AND

OF INVENTION: THERAPEUTIC METHODS

OF INVENTION: 335
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14-AUG-1997
                                                                                                                                                                                                                           Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 47;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "fusion protein composed of
enterokinase cleavable, His tagged
thioredoxin moiety and full length hTRT"
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RESULT 70
US-09-402-1818-600
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                                                                                                                                          Sequence 600, Application US/0940
Patent No. 6610839
GENERAL INFORMATION:
GENERAL TAPOLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                   Query Match 100.0%; Score 47; DB Best Local Similarity 100.0%; Pred. No. 35; Matches 6; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Say
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/912,951
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 1285 amino aci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 18-APR-1997
CLASSIFICATION: 435
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 25-APR-1997
CLASSIFICATION: 435
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 94111
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US 08/854,050
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Apple, Randolph REGISTRATION NUMBER: 30
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: Two Embarcadero Center, 8th Floor
           Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
                                                                                                                                                                                                                                                                                              32 WCGPCK 37
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SEQUENCES: 633
                                                                                     Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
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ID NO: 314:
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; NAME/KEY: Protein
; LOCATION: 1..1285
; OTHER INFORMATION: /note= "fusion protein composed of enterokinase cleavable, His tagged thioredoxin moiety and full length hTRT"
; SEQUENCE DESCRIPTION: SEQ ID NO: 600:
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Search completed: February 23, 2006, 00:43:50 Job time: 39.5 secs
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Best Local Similarity
Matches 6; Conserv
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FILING DATE: 09-MAY-1997

APPLICATION NUMBER: US 08/911,312

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/915,503

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: WS 08/915,503

FILING DATE: 01-OCT-1997

APPLICATION NUMBER: WD 067/17885

FILING DATE: 01-OCT-197

APPLICATION NUMBER: WD 07/17885

FILING DATE: 01-OCT-197

APPLICATION NUMBER: WD 07/17885

FILING DATE: 01-OCT-197

APPLICATION NUMBER: WD 07/17885

FILING DATE: 01-OCT-197

ATTORNEY/AGENT INFORMATION:

REGISTRATION NUMBER: 42,271

REFERENCE/DOCKET NUMBER: 015389-002620US

TELEPHONE: (415) 576-0300

NEORMATION INFORMATION:

TELEPHONE: (415) 576-0300
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APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/402,181B
FILING DATE: 29-Sep-1997
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: protein
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STRANDEDNESS: <Unknown>
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Sequence 4, Application US/10304287

Publication No. US20030083234A1

GENERAL INFORMATION:

APPLICANT: Walsman, David M.

APPLICANT: Walsman, David M.

APPLICANT: Kwon, Mijung

TITLE OF INVENTION: Anti-angiogenesis methods, compositions and uses therefor

FILE REFERENCE: MEGO-001

CURRENT APPLICATION NUMBER: US/10/304,287

CURRENT FILING DATE: 2001-11-26

PRIOR APPLICATION NUMBER: US 60/333,866

PRIOR FILING DATE: 2001-11-28

NUMBER OF SEQ ID NOS: 8

SOFTWARE: Microsoft Word

SEQ ID NO 4

LENGTH: 6

TYPE: PRT

ORGANISM: mammalian
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US-10-304-287-4
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Sequence 3, Application US/10660118A
Publication No. US20040131606A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum FILE REFERENCE: 2879-98
CURRENT APPLICATION NUMBER: US/10/660,118A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
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Best Local Similarity
Matches 6; Conserv
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100.0%; Pred. No. 1.7e+06;
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US-10-172-039A-459
US-10-210-028-459
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US-10-165-353A-459
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Sequence 4, Application US/09801260

Patent No. US20020034801A1

GENERAL INFORMATION:
APPLICANT: Curtis, Rory A.J.
ITILE OF INVENTION: 22105, A NOVEL HUMAN THIOREDOXIN FAMILY
TITLE OF INVENTION: MEMBER AND USES THEREOF
FILE REFERENCE: 10448-022001
CURRENT APPLICATION NUMBER: US/09/801,260
CURRENT FILING DATE: 2001-03-06
PRIOR APPLICATION NUMBER: US 60/187,447
PRIOR FILING DATE: 2000-03-07
NUMBER OF SEQ ID NOS: 5
SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 4
LENGTH: 52
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Sequence 4, Application No. US20040142897A1

GENERAL INFORMATION:
APPLICANT: Waisman, David M.
TITLE OF INVENTION: Compositions and Methods for Inhibiting Tum
FILE REFERENCE: ME03-009

CURRENT APPLICATION NUMBER: US/10/735,577

CURRENT FILING DATE: 2003-12-12

PRIOR APPLICATION NUMBER: US 60/433,140

PRIOR APPLICATION NUMBER: US 60/433,140

PRIOR APPLICATION NUMBER: US 60/433,140

SOFTWARE: Microsoft Word

SOFTWARE: Microsoft Word

SEQ ID NO. 4
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US-09-801-260-4
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US-10-735-577-4
                                                  ; OTHER INFORMATION: consensus sequence US-09-801-260-4
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SOFTWARE: Patentin version 3.1
SEQ ID NO 3
LENGTH: 6
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TYPE: PRT
ORGANISM: mammalian
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ORGANISM: Artificial
FEATURE:
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Pred. No.
 Score 47;
Pred. No.
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                 DB 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-425-115-351019
                                                                                                                                                                                                      APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 351019
                               Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE: 2002-05-14
Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 95
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 43
LENGTH: 52
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 351019, Application US/10425115 Publication No. US20040214272A1
                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                        OTHER INFORMATION: Clone ID: MRT4577_83299C.1.pep-10-425-115-351019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Bandaru, Rajasekhar
APPLICANT: Kapeller-Libermann, Rosana
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR FAMILY MEMBERS,
TITLE OF INVENTION: HUMAN THIOREDOXIN FAMILY MEMBERS, HUMAN LEUCINE-RIC
TITLE OF INVENTION: REPEAT FAMILY MEMBERS, AND HUMAN RINGFINGER FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Alexandra Glucksmann, Maria
APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: M. Galvin, Katherine
                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: La Rosa, Thomas J. APPLICANT: Kovalic, David K.
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                                                                                                                                                    LENGTH: 82
TYPE: PRT
ORGANISM: Zea mays
                                                                                                                                            FEATURE:
                               Local Similarity
nes 6; Conserv
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                                   Conservative
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                               100.0%; Score 47; DB 4; Length 82; 100.0%; Pred. No. 10; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 47; DB 4; Length 52; 100.0%; Pred. No. 7.1;
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APPLICANT: Ghosh, Soumitra S.
APPLICANT: Fahy, Boin D.
APPLICANT: Zhang, Bing
APPLICANT: Zhang, Bing
APPLICANT: Zhang, Bradford W.
APPLICANT: Glbson, Bradford W.
APPLICANT: Taylor, Steven W.
APPLICANT: Warnock, Dale E.
APPLICANT: Warnock, Dale E.
TITLE OF INVENTION: TARGETS FOR THERAPEUTIC INTERVENTION
TITLE OF INVENTION: TARGETS IN THE MITOCHONDRIAL PROTEOME
FILE REFERENCE: 660088.465
FILE REFERENCE: 60088.465
FILE REFERENCE: 60088.465
CURRENT APPLICATION NUMBER: US/10/408,765A
CURRENT APPLICATION STANDARD CONTROL FILING DATE: 2003-04-04
NUMBER OF EGG ID NOS: 3077
SOPTWARE: PastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53223)B

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 285684

SEQ ID NO 280571
LENGTH: 90

TYPE: ppm
                                                                                                                                                              ; OTHER INFORMATION: Clone ID: PAT_MRT3847_95379C.1.pep US-10-424-599-280571
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US-10-424-599-280571
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-408-765A-312
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US-10-408-765A-312
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Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                             NAME/KEY: unsure
LOCATION: (1)...(90)
OTHER INFORMATION: unsure at all Xaa locations
                                                                                                                                                                                                                                                                                                         ORGANISM: Glycine max
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                                                                           6; Conserv
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WCGPCK 18
                                       WCGPCK 6
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o. US20040101874A1
                                                                           100.0%; Score 47; DB ilarity 100.0%; Pred. No. 11; Conservative 0; Mismatches
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                                                                                                                     DB 4; Length 90;
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FILE REPERENCE: 38814 3518
CURRENT APPLICATION NUMBER: US/10/032,2018
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 102
LENGTH: 102
TYPE: PRT
ORGANISM: Chlamydia muridarum
US-10-032-2018-102
RESULT 11
US-10-032-201B-104
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US-10-032-201B-102
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; ORGANISM: Thiocapsa roseopersicina
US-10-032-201B-143
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US-10-032-201B-143
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 143
LENGTH: 91
                                                                                                                                                                        Query Match 100.0%; Score 47; DB 4; Length 102; Best Local Similarity 100.0%; Pred. No. 12; Matches 6; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Briggs Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
APPLICANT: Briggs, Steven
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APPLICANT: Deckers, Harm
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                                                                                        27 WCGPCK 32
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Dalmia, Bipin Kumar
Del Val, Greg
Zaplachinski, Steve
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Heifetz, Peter Bernard
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Heifetz, Peter Bernard
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RESULT 13

US-10-032-201B-126

; Sequence 126, Application US/10032201B

; Publication No. US20030167524A1

; GENERAL INFORMATION:
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US-10-032-201B-105
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NUMBER OF SEQ ID NOS: 313

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 105

LENGTH: 102

TYPE: PRT

ORGANISM: Chlamydia trachomatis
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NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 104
                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.
Matches 6; Conservative
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Best Local Similarity 100
Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Van Rodijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bernard
APPLICANT: Heifetz, Peter Bernard
APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Zaplachinski, Steve
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
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APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
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APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bernard
APPLICANT: Briggs, Steven
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ORGANISM: Psittaci
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Dalmia, Bipin Kumar
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100.0%; Pred. No.
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RESULT 15
US-10-501-282-3126
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                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 128
LENGTH: 102
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 126
LENGTH: 102
                                                     Sequence 3126, Application US/10501282 Publication No. US20050203280A1 GENERAL INFORMATION:
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Publication No. US20030167524A1
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Best Local Similarity
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APPLICANT:
APPLICANT:
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Best Local Similarity
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APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
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APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS,
TITLE OF INVENTION: COMPOSITIONS
APPLICANT: MCMICHAEL, JOHN CALHOUN
APPLICANT: ZAGURSKY, ROBERT JOHN
APPLICANT: RUSSELL, DAVID PARRISH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION
TITLE OF INVENTION: COMPOSITIONS
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Zaplachinski, Steve
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Dalmia, Bipin Kumar
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Dalmia, Bipin Kumar
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100.0%; Pred. No.
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Pred. No.
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RESULT 17
US-10-032-585-7785
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US-10-032-201B-96
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SEQ ID NO 96
SERVIE 103
TYPE: PRT
ORGANISM: Bacillus subtilis
                                                               GENERAL INFORMATION:
APPLICANT: Terry, Roemer D.
                                                                                                              Sequence 7785, Application US/10032585 Publication No. US20030180953A1
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Best Local Similarity
Matches 6; Conserv
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APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bernard
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SEQ ID NO 3126
LENGTH: 102
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APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
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Best Local
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TITLE OF INVENTION: ALLOIOCOCCUS OTITIDIS OPEN READING FRAMES (ORFS) ENCODING
TITLE OF INVENTION: POLYPEPTIDE ANTIGENS, IMMUNOGENIC COMPOSITIONS AND USES THEREOF
FILE REFERENCE: AM100780 L2
CURRENT APPLICATION NUMBER: US/10/501,282
CURRENT FILING DATE: 2004-07-09
PRIOR APPLICATION NUMBER: 60/333,777
PRIOR EPILING DATE: 2001-11-29
PRIOR APPLICATION NUMBER: 60/426,742
PRIOR APPLICATION NUMBER: 60/426,742
PRIOR APPLICATION NUMBER: FCT/US02/36123
PRIOR APPLICATION NUMBER: PCT/US02/36123
PRIOR APPLICATION NUMBER: PCT/US02/36123
PRIOR FILING DATE: 2002-11-25
PRIOR FILING DATE: 2002-11-25
PRIOR FILING DATE: 2002-11-25
PRIOR APPLICATION NUMBER: PCT/US02/36123
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CURRENT APPLICATION UNMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
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TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
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Dalmia, Bipin Kumar
Del Val, Greg
Caplachinski, Steve
                  Bo, Jiang
Charles, Boone
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Wo. US20030167524A1
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Query Match
Best Local Similarity
Watches 6; Conserve
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Best Local Similarity
Thes 6; Conserv
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US-10-032-585-7785
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CURRENT FILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 8000
SOFTWARE: PatentIn version 3.1
SEQ ID NO 7785
LENGTH: 103
                                                                                                                                                                                                                                                                                                              Sequence 270, Application US/10451467A Publication No. US20040161840A1 GENERAL INFORMATION:
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SEQ ID NO 7
LENGTH: 103
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                                       CURRENT APPLICATION NUMBER: US/10/451,467A CURRENT FILING DATE: 2003-06-19 PRIOR APPLICATION NUMBER: EP 00870318.3 PRIOR FILING DATE: 2000-12-22 PRIOR APPLICATION NUMBER: EP 01870002.1 PRIOR FILING DATE: 2001-01-04
                                                                                                                                                                                                                                APPLICANT: CONTRERAS, ROLAND HENRI
APPLICANT: EBERHARDT, INES
APPLICANT: LUYTEN, WALTER HERMAN MARIA LOUIS
APPLICANT: REEKMANS, RIEKA JOSEPHINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/660,118A CURRENT FILING DATE: 2003-09-10 PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or
FILE REFERENCE: 2879-98
  PRIOR APPLICATION NUMBER: EP 01870003.9 PRIOR FILING DATE: 2001-01-09
                                                                                                                                                                       FILE REFERENCE: JAB-1667
                                                                                                                                                                                        TITLE OF INVENTION: BAX-RESPONSIVE GENES FOR DRUG TARGET IDENTIFICATION TITLE OF INVENTION: YEAST AND FUNGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 15
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Pred. No. 12;
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CURRENT FILING DATE: 2003-06-19
PRIOR APPLICATION NUMBER: EP 00870318.3
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: EP 01870002.1
PRIOR APPLICATION NUMBER: EP 01870002.1
PRIOR FILING DATE: 2001-01-04
PRIOR FILING DATE: 2001-01-09
PRIOR FILING DATE: 2001-01-09
NUMBER OF SEQ ID NOS: 732
SOFTWARE: Patentin version 3.1
SEQ ID NO 450
LENGTH: 103
TYPE: PRT
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Best Local Similarity
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                                                                                                                                                                Publication No. US20040161840A1
GENERAL INFORMATION:
APPLICANT: CONTRERAS, ROLAND HENRI
APPLICANT: EBERHARDT, INES
APPLICANT: LUYTEN, WALTER HERMAN MARIA LOUIS
APPLICANT: REEKMANS, RIEKA JOSEPHINA
TITLE OF INVENTION: BAX-RESPONSIVE GENES FOR DRUG TARGET IDENTIFICATION IN
TITLE OF INVENTION: YEAST AND FUNGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best
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                                    FILE REFERENCE: JAB-1667
CURRENT APPLICATION NUMBER: US/10/451,467A
CURRENT FILING DATE: 2003-06-19
PRIOR APPLICATION NUMBER: EP 00870318.3
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: EP 01870002.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: CONTRERAS, ROLAND HENRI APPLICANT: EBERHARDT, INES
PRIOR FILING DATE: 2001-01-04
PRIOR APPLICATION NUMBER: EP 01870003.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Candida albicans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Saccharomyces cerevisiae
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nes 6; Conservativo
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100.0%; Pred. No.
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Pred. No. 12;
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Gaps

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Length 103; Indels

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SOFTWARE: PatentIn version
SEQ ID NO 270
LENGTH: 103
APPLICANT: EBERHARDT, INES
APPLICANT: LUYTEN, WALTER HERMAN MARIA LOUIS
APPLICANT: REEKMANS, RIEKA JOSEPHINA
TITLE OF INVENTION: BAX-RESPONSIVE GENES FOR DRUG TARGET IDENTIFICATION IN
TITLE OF INVENTION: YEAST AND FUNGI
FILE REFERENCE: JAB-1667
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NUMBER OF SEQ ID NOS: 732

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CURRENT APPLICATION NUMBER: US/09/954,342
CURRENT FILING DATE: 2001-09-17
PRIOR APPLICATION NUMBER: 60/233,382
PRIOR FILLING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: 60/240,498
PRIOR PILLING DATE: 2000-10-13
PRIOR FILLING DATE: 2000-10-13
PRIOR FILLING DATE: 2001-01-08
PRIOR FILLING DATE: 2001-01-12
PRIOR APPLICATION NUMBER: 60/264,794
PRIOR FILLING DATE: 2001-01-12
PRIOR APPLICATION NUMBER: 60/238,398
PRIOR FILLING DATE: 2001-0-06
PRIOR APPLICATION NUMBER: 60/232,675
PRIOR APPLICATION NUMBER: 60/232,675
PRIOR FILLING DATE: 2000-09-15
PRIOR PILLING DATE: 2001-03-09
PRIOR FILLING DATE: 2001-03-09
PRIOR FILLING DATE: 2001-03-09
PRIOR FILLING DATE: 2000-09-15
PRIOR PILLING DATE: 2000-09-15
PRIOR APPLICATION NUMBER: 60/233,801
PRIOR PILLING DATE: 2000-09-15
PRIOR APPLICATION NUMBER: 60/233,601
PRIOR PILLING DATE: 2000-09-15
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SEQ ID NO 614
LENGTH: 103
TYPE: PRT
ORGANISM: Candida albicans
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Best Local Similarity
Matches 6; Conserv
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APPLICANT: SPYTEK, KIMBERLY ANN
APPLICANT: TAUPIER, RAYMOND J.
APPLICANT: VERNET, CORINE A.
APPLICANT: COLMAN, STEVEN D.
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APPLICANT:
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APPLICANT:
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                                      APPLICATION NUMBER: 60/:
PTLING DATE: 2000-09-20
                  PILING DATE: 2000-09-20
APPLICATION NUMBER: 60/233,402
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PADIGARU, MUKAL...

GERLACH, VALERIE L.

MACIDOUGALL, JOHN R.

MACIDOUGALL, JOHN R.

TOTALE
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GUNTHER, ERIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RASTELLI, LUCA
ZERHUSEN, BRYA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ELLERMAN, KAREN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TCHERNEV, VELIZAR T. MALYANKAR, URIEL M. SHENOY, SURESH
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2000-09-18
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                                                             60/233,960
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Pred. No.
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12;
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APPLICANT: EASTELLI, LUCA
APPLICANT: ZERHUSEN, BRYAN
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES AND POLYPEPTIDES ENCODED THEREBY
FILE REFERENCE: 21402-114
CURRENT FILING DATE: 2001-09-17
CURRENT FILING DATE: 2001-09-17
PRIOR APPLICATION NUMBER: 60/233,382
PRIOR APPLICATION NUMBER: 60/240,498
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: 60/240,498
PRIOR FILING DATE: 2001-01-01
PRIOR APPLICATION NUMBER: 60/260,284
PRIOR FILING DATE: 2001-01-08
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/260,973
PRIOR FILING DATE: 2001-01-12
PRIOR APPLICATION NUMBER: 60/264,794
PRIOR FILING DATE: 2001-01-29
PRIOR APPLICATION NUMBER: 60/238,398
PRIOR FILING DATE: 2000-10-61
PRIOR APPLICATION NUMBER: 60/232,675
PRIOR APPLICATION NUMBER: 60/232,675
PRIOR PILING DATE: 2000-09-15
PRIOR PILING DATE: 2000-09-15
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PRIOR APPLICATION NUMBER: 60/232,675
PRIOR PILING DATE: 2000-09-15
PRIOR PELICATION NUMBER: 60/274,862
PRIOR PILING DATE: 2001-03-09
PRIOR PILING DATE: 2001-03-09
PRIOR APPLICATION NUMBER: 60/233,801
PRIOR APPLICATION NUMBER: 60/232,676
PRIOR APPLICATION NUMBER: 60/232,676
PRIOR APPLICATION NUMBER: 60/232,960
PRIOR APPLICATION NUMBER: 60/233,960
PRIOR PILING DATE: 2000-09-20
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Best Local S
Matches 6
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Publication No. US20030170838
GENERAL INFORMATION:
APPLICANT: MISHAA, VISHNU S.
APPLICANT: SPYTEK, KIMBERLY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 62
LENGTH: 104
TYPE: PRT
ORGANISM: Ovis aries
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PRIOR FILING DATE: 2000-09-19
PRIOR, APPLICATION NUMBER: 60/232,679
PRIOR FILING DATE: 2000-09-15
NUMBER OF SEQ ID NOS: 104
SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GUNTHER, ERIK
ELLERMAN, KAREN
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GERLACH, VALERIE L.
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No. US20030170838A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LINDA
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100.0%; Pred. No.
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                                                                                                                                        PRIOR APPLICATION NUMBER: 60/233,382
PRIOR FILLING DATE: 2000-09-18
PRIOR PELICATION NUMBER: 60/240,498
PRIOR FILLING DATE: 2000-10-13
PRIOR APPLICATION NUMBER: 60/260,284
PRIOR APPLICATION NUMBER: 60/260,973
PRIOR APPLICATION NUMBER: 60/260,973
PRIOR APPLICATION NUMBER: 60/264,794
PRIOR APPLICATION NUMBER: 60/264,794
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 63
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Publication No. US20030170838A1
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APPLICANT: SPYTEK, KIMBERLY ANN
APPLICANT: TAUPIER, RAYMOND J.
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                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES AND POLYPEPTIDES ENCODED THEREBY FILE REFERENCE: 21402-114 CURRENT APPLICATION NUMBER: US/09/954,342 CURRENT FILING DATE: 2001-09-17
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                                                 FILING DATE: 2001-01-29
APPLICATION NUMBER: 60/238,398
FILING DATE: 2000-10-06
APPLICATION NUMBER: 60/232,675
FILING DATE: 2000-09-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/233,521
FILING DATE: 2000-09-19
APPLICATION NUMBER: 60/233,522
FILING DATE: 2000-09-19
               APPLICATION NUMBER: 60/274,862
FILING DATE: 2001-03-09
APPLICATION NUMBER: 60/233,801
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6; Conserva
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GERLACH, VALERIE L.
MACDOUGALL, JOHN R.
SMITHSON, GLENNDA
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PEYMAN, JOHN
STONE, DAVID
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COLMAN, STEVEN D.
GORMAN, LINDA
                                                                                                                                                                                                                                                                                                                                                                                   ZERHUSEN, BRYAN
                                                                                                                                                                                                                                                                                                                                                                                                  RASTELLI, LUCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           GUNTHER, ERIK
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RESULT 25
US-09-954-342-65
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APPLICANT: ZERHUSEN, BYYAN
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES AN
FILE REFERENCE: 21402-114
CURRENT APPLICATION NUMBER: US/09/954,342
CURRENT FILING DATE: 2001-09-17
PRIOR APPLICATION NUMBER: 60/233,382
PRIOR FILING DATE: 2000-09-18
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: 60/240,498
PRIOR FILING DATE: 2000-10-13
PRIOR APPLICATION NUMBER: 60/260,284
PRIOR APPLICATION NUMBER: 60/260,284
PRIOR FILING DATE: 2001-01-08
PRIOR FILING DATE: 2001-01-08
PRIOR FILING DATE: 2001-01-08
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SEQ ID NO 64
LENGTH: 104
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PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: 60/233,521
PRIOR FILING DATE: 2000-09-19
PRIOR APPLICATION NUMBER: 60/233,522
PRIOR APPLICATION NUMBER: 60/233,522
PRIOR APPLICATION NUMBER: 60/232,679
PRIOR APPLICATION NUMBER: 60/232,679
PRIOR APPLICATION NUMBER: 60/260,973
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/264,794
PRIOR FILING DATE: 2001-01-29
PRIOR APPLICATION NUMBER: 60/238,398
PRIOR FILING DATE: 2000-10-06
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 60/232,675
PRIOR FILING DATE: 2000-09-15
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Best Local Similarity
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APPLICANT: SPYTEK, KIMBERLY ANN
APPLICANT: TAUPIER, RAYMOND J.
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NUMBER OF SEQ ID NOS: 104
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PRIOR FILING DATE: 2000-09-15
PRIOR APPLICATION NUMBER: 60/233,960
PRIOR FILING DATE: 2000-09-20
PRIOR FILING DATE: 2000-09-20
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GERLACH, VALERIE L.
GONYGALL, JOHN R.
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STONE, DAVID
GUNTHER, ERIK
ELLERMAN, KAREN
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MACDOUGALL, UC...
MACDOUGALL, UC...
GLENNDA
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FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 146
SEQ ID NO 146
LENGTH: 104
TYPE: PRT
ORGANISM: Bos taurus
US-10-032-201B-146
RESULT 27
US-10-032-201B-150
; Sequence 150, Application US/10032201B
; Publication No. US20030167524A1
; GENERAL INFORMATION:
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Best Local Similarity
Marches 6; Conserve
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US-10-032-201B-146
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Best Local Similarity
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TITLE OF INVENTION: METHODS FOR THE PRODUCTION
TITLE OF INVENTION: COMPOSITIONS
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APPLICANT: Deckers, Harm
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TYPE: PRT
ORGANISM: Rattus norvegicus
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Dalmia, Bipin Kumar
Del Val, Greg
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Heifetz, Peter Bernard
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Pred. No. 12;
                                                                                                                                                                                            Score 47; DB Pred. No. 12;
                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                            Length 104;
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APPLICANT: Van Rooijen, Gijs APPLICANT: Deckers, Harm APPLICANT: Heifetz, Peter Bernard

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RESULT 29

US-10-032-201B-154

; Sequence 154, Application US/10032201B

; Publication No. US20030167524A1

; GENERAL INFORMATION:
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US-10-032-201B-152
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 150
                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 6; Conserv
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SEQ ID NO 152
LENGTH: 104
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Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
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APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter B
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APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION
TITLE OF INVENTION: COMPOSITIONS
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APPLICANT:
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TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
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ORGANISM: Gallus gallus
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                                                                                                                                    30 WCGPCK 35
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Heifetz, Peter Bernard
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Dalmia, Bipin Kumar
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Del Val, Greg
Zaplachinski, Steve
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milarity 100.0%;
Conservative C
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Query Match
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US-10-032-201B-158
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                                                 Sequence 158, Application US/10032201B
Publication No. US20030167524A1
GENERAL INFORMATION:
APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bernard
                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 155
LENGTH: 104
TYPE: PRT
ORGANISM: Mus musculus
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 154
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APPLICANT:
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Best Local
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APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
TITLE REFERENCE: 38814 351B
                 APPLICANT:
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APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
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APPLICANT: Deckers, Harm
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TYPE: PRT
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Local Similarity 100.0%;
hes 6; Conservative
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Heifetz, Peter Bernard
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Dalmia, Bipin Kumar
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Dalmia, Bipin Kumar
                 Briggs, Steven
Dalmia, Bipin Kumar
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Greg
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APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
ITITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND
FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 158
LENGTH: 104
TYPE: PRT
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APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bernard
APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
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SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 159
LENGTH: 104
TYPE: PRT
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Matches
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Best Local Similarity
Matches 6; Conserv
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                 APPLICANT:
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APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
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CURRENT FILING DATE: 2001-12-19
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nes 6; Conser
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Heifetz, Peter Bernard
                                                               Briggs, Steven
Dalmia, Bipin Kumar
Moloney, Maurice
ENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS,
                             Del Val, Greg
Zaplachinski, Steve
                                                                                                                                                                Application US/10032201B
lo. US20030167524A1
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o. US20030167524A1
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100.0%; Pred. No.
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Pred. No.
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RESULT 35
US-10-424-599-282588
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US-10-032-201B-161
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Sequence 282588, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Cao Yongwei
APPLICANT: Cao Yongwei
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 282588
LENGTH: 104
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 276
LENGTH: 104
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Publication No. US20030167524A1
GENERAL INFORMATION:
APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
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Matches
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 161
LENGTH: 104
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APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Staphylococcus aureus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
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100.0%; Pred. No.
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Pred. No.
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US-09-954-342-61
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SEQ ID NO 188
LENGTH: 104
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APPLICANT: MISHRA, VISHNU S.
APPLICANT: SPYTEK, KIMBERLY
APPLICANT: TAUPIER, RAYMOND
                                                                                                                                                                                                                                                                                                    Sequence 61, Application US/09954342 Publication No. US20030170838A1
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: REEKMANS, RIEKA JOSEPHINA
TITLE OF INVENTION: BAX-RESPONSIVE GENES FOR DRUG TARGET IDENTIFICATION:
TITLE OF INVENTION: YEAST AND FUNGI
FILE REFERENCE: JAB-1667
CURRENT APPLICATION NUMBER: US/10/451,467A
CURRENT FILING DATE: 2003-06-19
PRIOR APPLICATION NUMBER: EP 00870318.3
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: EP 01870002.1
PRIOR FILING DATE: 2001-01-04
PRIOR PILING DATE: 2001-01-04
PRIOR PILING DATE: 2001-01-09
PRIOR FILING DATE: 2001-01-09
PRIOR FILING DATE: 2001-01-09
PRIOR FILING DATE: 2001-01-09
PRIOR FILING DATE: 2001-01-09
                                                                                     APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: CONTRERAS, ROLAND HENRI
APPLICANT: EBERHARDT, INES
APPLICANT: LUYTEN, WALTER HERMAN MARIA LOUIS
                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 732
                                                      APPLICANT
                                                                                                                         APPLICANT:
                                                                                                                                                            APPLICANT:
                                   APPLICANT
                                                                                                                                          APPLICANT:
                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Saccharomyces cerevisiae
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Clone ID: PAT_MRT3847_97199C.1.pep
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    54 WCGPCK 59
                                                                                                                                                                                                                                                                                                                                                                                                             30 WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                1 WCGPCK 6
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                                                         PADIGARU, MUKALLARIE L.
GERLACH, VALERIE L.
TOTICALL, JOHN R.
                                                                                                                                                                                                                             SPYTEK, KIMBERLY ANN TAUPIER, RAYMOND J.
                                                                                                                                                                         VERNET, CORINE A.
COLMAN, STEVEN D.
GORMAN, LINDA
 STONE,
                                                  MACDOUGALL, JOHN I
SMITHSON, GLENNDA
                                                                                                                         SHENOY, SURESH
                                                                                                                                        TCHERNEV, VELIZAR T. MALYANKAR, URIEL M.
                PEYMAN,
                               MILLET, ISABELLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/10451467A
o. US20040161840A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 47; DB 100.0%; Pred. No. 12;
                                                                                                      MURALIDHARA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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RESULT 38
US-10-171-311-226
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Publication No. US20030087270A1
GENERAL INFORMATION:
APPLICANT: Schlegel, Robert
APPLICANT: Chen, Yan
APPLICANT: Zhao, Xumei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver.
SEQ ID NO 61
LENGTH: 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Equus caballus -09-954-342-61
              APPLICANT: Monahan, John
APPLICANT: Kamatkar, Shubhangi
APPLICANT: Glatt, Karen
APPLICANT: Glatt, Karen
APPLICANT: Gannavarapu, Manjula
APPLICANT: Hoersh, Sebastian
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS
TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND
TITLE OF INVENTION: OF CERVICAL CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 2001-09-17
PRIOR APPLICATION NUMBER: 60/233,382
PRIOR FILING DATE: 2000-09-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE: 2001-09-17
FILE
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NUMBER OF SEQ ID NOS: 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/232,679 PRIOR FILING DATE: 2000-09-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/260,284
PRIOR FILING DATE: 2001-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR EILING DATE: 2000-10-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/260,973
PRIOR FILING DATE: 2001-01-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES AND POLYPEPTIDES ENCODED THEREBY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
mes 6; Conserv
REFERENCE: MRI-035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/264,794 FILING DATE: 2001-01-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 60/233,522
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Pred. No.
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Mismatches
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CURRENT FILING DATE: 2002-06-12
PRIOR APPLICATION NUMBER: US 60/298,159
PRIOR FILING DATE: 2001-06-13
PRIOR APPLICATION NUMBER: US 60/298,155
PRIOR FILING DATE: 2001-06-13
PRIOR FILING DATE: 2001-06-13
PRIOR APPLICATION NUMBER: US 60/335,936
PRIOR APPLICATION NUMBER: US 60/335,936
PRIOR APPLICATION NUMBER: US 60/335,936
PRIOR FILING DATE: 2001-11-14
                                                                      APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Ber
APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kuma
APPLICANT: Del Val, Greg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 291, Application US/10316253 Publication No. US20030162706A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 238
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 291
LENGTH: 105
                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                            Sequence 47, Appropriation No.
                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/316,253
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 60/355,295
PRIOR FILING DATE: 2002-02-08
NUMBER OF SEQ ID NOS: 308
SOFTWARE: PatentIn version 3.1
                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Greis, Kenneth
TITLE OF INVENTION: Angiogenesis Modulating
FILE REFERENCE: 8865M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: The Procter & Gamble Company
APPLICANT: Peters, Kevin
APPLICANT: Thompson, Larry
APPLICANT: Wang, Feng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 105
TYPE: PRT
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mes 6; Conserv
OF INVENTION:
                                                                                                                                                                                                                                                                                                                               31 WCGPCK 36
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                                                                                                                                                                                                                                                                                                                                                                   1 WCGPCK 6
                    INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                 6;
Moloney, Maurice vention: Methods for the production of multimeric proteins, and related vention: compositions
                                                                                                                                Deckers, Harm
Heifetz, Peter Bernard
                                                                                                                                                                                                            Application US/10032201B
vo. US20030167524A1
                                                        Del Val, Greg
Zaplachinski, Steve
                                                                                          Briggs, Steven
Dalmia, Bipin Kumar
                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 47;
100.0%; Pred. No.
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFFWARE: FASLSEQ for Windows Version 4.0
SEQ ID NO 47
LENGTH: 105
TYPE: PRT
ORCANISM: Homo Sapien
US-10-032-201B-47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/032,201B; CURRENT FILING DATE: 2001-12-19; NUMBER OF SEQ ID NOS: 313; SOFTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 106; LENGTH: 105

TYPE: DOWN
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US-10-424-599-237784
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US-10-032-201B-106
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APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT APPLICATION ONMER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 237784
LENGTH: 105
                                                                                                                                                                                                                              Sequence 237784, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 106, Application US/10032201B Publication No. US20030167524A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
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Del Val, Greg
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Dalmia, Bipin Kumar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Heifetz, Peter Bernard
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Pred. No.
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 127
LENGTH: 105
TYPE: PRT
ORGANISM: Mus musculus
PUBLICATION INFORMATION:
DATABASE EXTRY DATE: 2002-12-25
US-10-351-891-127
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US-10-351-891-127
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                                                                                                                                                                                                                                                                                                                                               US-10-408-765A-2019
SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 2019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 127, Application US/10351891
Publication No. US20040048311A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local
Matches
                             APPLICANT: Gibson, Bradford W.
APPLICANT: Taylor, Steven W.
APPLICANT: Taylor, Steven W.
APPLICANT: Glenn, Gary M.
APPLICANT: Warnock, Dale E.
TITLE OF INVENTION: TARGETS FOR THERAPEUTIC INTERVENTION
TITLE OF INVENTION: IDENTIFIED IN THE MITOCHONDRIAL PROTEOME
FILE REPERENCE: 660088.465
CURRENT APPLICATION NUMBER: US/10/408,765A
CURRENT FILING DATE: 2003-04-04
NUMBER OF SEQ ID NOS: 3077
                                                                                                                                                                                                                                                                                                       Sequence 2019, Application US/10408765A Publication No. US20040101874A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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TITLE OF INVENTION: USE OF COLLECTIONS OF BINDING
TITLE OF INVENTION: APPLICATIONS
FILE REFERENCE: 25885-1753
CURRENT APPLICATION NUMBER: US/10/351,891
CURRENT FILING DATE: 2003-01-24
PRIOR APPLICATION NUMBER: US 60/352,011
PRIOR PILING DATE: 2002-01-24
NUMBER OF SEO ID NOS: 140
NUMBER OF SEO ID NOS: 140
                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: DANA AULT-RICHE APPLICANT: PAUL D. KASSNE
                                                                                                                                                                                                                                                                 APPLICANT: Ghosh, Soumitra S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: unsure
LOCATION: (1)..(105)
OTHER INFORMATION: u
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 RESULT 45
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 ; ORGANISM: Homo sapiens US-10-408-765A-2019
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 US-10-660-118A-9
 US-10-660-118A-8
 US-10-660-118A-9
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Matches 6
 Sequence 9, Application US/10660118A
Publication No. US20040131606A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum
FILE REFERENCE: 2879-98
FILE REFERENCE: 2879-98
 Sequence 8, Application US/10660118A Publication No. US20040131606A1 GENERAL INFORMATION:
APPLICANT: White, Carl W.
 NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.1
SEQ ID NO 9
 Matches
 Query Match
 SEQ ID NO 8
 CURRENT APPLICATION NUMBER: US/10/660,118A CURRENT FILING DATE: 2003-09-10 PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10 PRIOR APPLICATION NUMBER: 60/462,082 PRIOR FILING DATE: 2003-04-11
 TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum FILE REFERENCE: 2879-98

CURRENT APPLICATION NUMBER: US/10/660,118A

CURRENT FILING DATE: 2003-09-10

PRIOR APPLICATION NUMBER: 60/409,960

PRIOR FILING DATE: 2002-09-10

PRIOR APPLICATION NUMBER: 60/462,082
 NUMBER OF SEQ ID NOS: 15
 SOFTWARE: PatentIn version 3.1
 PRIOR FILING DATE:
 TYPE: PRT
ORGANISM: Mus musculus
 TYPE: PRT
ORGANISM: Gallus gallus
 LENGTH: 105
 ENGTH: 105
 Local Similarity
 31
 31 WCGPCK 36
 31
 1 WCGPCK 6
 1 WCGPCK 6
 1 WCGPCK 6
 Similarity 6; Conserv
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WCGPCK 36
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 2003-04-11
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 Score 47;
Pred. No.
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 Score 47; DB 4;
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 RESULT 49
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 US-10-660-118A-10
 US-10-660-118A-11
 US-10-660-118A-11
 RESULT 48
 Sequence 12, Application US/10660118A
Publication No. US20040131606A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum
FILE REFERENCE: 2879-98
 Sequence 10, Application US/10660118A
publication No. US20040131606A1
publication No. US20040131606A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum
FILE REFERENCE: 2879-98
CURRENT APPLICATION NUMBER: US/10/660,118A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/462,082
prior APPLICATION NUMBER: 60/462,082
 Sequence 11, Application US/10660118A Publication No. US20040131606A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.1 SEQ ID NO 11
 Matches
 SEQ ID NO 10
 Query Match
Best Local Similarity
 Matches
 Query Match
 CURRENT APPLICATION NUMBER: US/10/660,118A CURRENT FILING DATE: 2003-09-10 PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10 PRIOR APPLICATION NUMBER: 60/462,082
 CURRENT APPLICATION NUMBER: US/10/660,118A
 PRIOR FILING DATE: 2003-
 SOFTWARE: PatentIn version 3.1
 NUMBER OF SEQ ID NOS:
 PRIOR FILING DATE: 2003-04-11
 LENGTH: 105
TYPE: PRT
ORGANISM: Rattus norvegicus
 ORGANISM: Bos taurus
 TYPE: PRT
 LENGTH:
 Local Similarity
 31 WCGPCK 36
 105
 31 WCGPCK 36
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 1 WCGPCK 6
 1 WCGPCK 6
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 RESULT 50
US-10-451-467A-684
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 ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-660-118A-12
 US-10-699-113-32
 US-10-451-467A-684
 SOFTWARE: PatentIn
SEQ ID NO 684
LENGTH: 105
 Sequence 684, Application US/10451467A Publication No. US20040161840A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.1
SEQ ID NO 12
LENGTH: 105
 Sequence 32, Application US/10699113 Publication No. US20040241748A1 GENERAL INFORMATION:
 Matches
 Matches
 Query Match
Best Local Similarity
 Query Match
Best Local Similarity
 APPLICANT: Ault-Riche, Dana
APPLICANT: Kumble, Krishnanand
APPLICANT: Schulz, Rainer
APPLICANT: Schulz, Reiner
TITLE OF INVENTION: Self-Assembling Arrays and Uses Thereof
FILE REFERENCE: 25885-1755
CURRENT APPLICATION NUMBER: US/10/699,113
CURRENT FILING DATE: 2003-10-30
 PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: EP 01870002.1
PRIOR FILING DATE: 2001-01-04
 CURRENT APPLICATION NUMBER: US/10/451,467A
CURRENT FILING DATE: 2003-06-19
PRIOR APPLICATION NUMBER: EP 00870318.3
 APPLICANT: REEKMANS, RIEKA JOSEPHINA
TITLE OF INVENTION: BAX-RESPONSIVE GENES FOR DRUG TARGET IDENTIFICATION IN
TITLE OF INVENTION: YEAST AND FUNGI
FILE REFERENCE: JAB-1667
 PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 15
 CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/409,960
 NUMBER OF SEQ ID NOS: 732
 PRIOR FILING DATE: 2001-01-09
 PRIOR APPLICATION NUMBER: EP 01870003.9
 APPLICANT:
 APPLICANT: CONTRERAS, ROLAND HENRI
 PRIOR FILING DATE:
PRIOR APPLICATION NUMBER: 60/446,687
 TYPE: PRT
ORGANISM: Homo sapiens
 31 WCGPCK 36
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
 in 100.0%; Score 47; Similarity 100.0%; Pred. No. 6; Conservative 0; Mismatch
 6;
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LUYTEN, WALTER HERMAN MARIA LOUIS
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 2002-09-10
 100.0%; Score 47; DB 4; 100.0%; Pred. No. 12;
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 US-10-970-370-2
; ORGANISM: Macaca mulatta US-10-970-370-6
 RESULT 53
US-10-970-370-6
 US-10-970-370-2
 US-10-699-113-32
 Query Match 100.0%; Score 47; DB Best Local Similarity 100.0%; Pred. No. 12; Matches 6; Conservative 0; Mismatches
 NUMBER OF SEQ ID NOS: 9
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
LENGIH: 105
 PRIOR FILING DATE: 2003-02-10
NUMBER OF SEQ ID NOS: 948
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 32
 SOFTWARE: PatentIn Ver. SEQ ID NO 6
 GENERAL INFORMATION:
 Sequence 2, Application US/10970370 Publication No. US20050208037A1
 Matches
 Query Match
Best Local S
 CURRENT APPLICATION NUMBER: US/10/970,370
CURRENT FILING DATE: 2004-10-21
PRIOR APPLICATION NUMBER: 60/513,134
PRIOR FILING DATE: 2003-10-21
NUMBER: OF SEQ ID NOS: 9
 GENERAL INFORMATION:
 Sequence 6, Application US/10970370 Publication No. US20050208037A1
 CURRENT APPLICATION NUMBER: US/10/970,370
CURRENT FILLING DATE: 2004-10-21
PRIOR APPLICATION NUMBER: 60/513,134
PRIOR FILING DATE: 2003-10-21
 TITLE OF INVENTION: THIOREDOXIN INCREASES REDOX-CYCLING OF ITITLE OF INVENTION: AGENTS THEREBY SENSITIZES CANCER CELLS FILE REFERENCE: UTSN:025US
 TITLE OF INVENTION: THIOREDOXIN INCREASES REDOX-CYCLING OF ANTICANCER TITLE OF INVENTION: AGENTS THEREBY SENSITIZES CANCER CELLS TO APOPTOSIS FILE REFERENCE: UTSN:025US
 LENGTH: 105
TYPE: PRT
ORGANISM: Mus musculus
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank BAA04881
DATABASE ENTRY DATE: 2002-12-25
 APPLICANT: DASHNAMOORTHY, RAVI APPLICANT: DAS, KUMUDA C.
 APPLICANT: DASHNAMOORTHY, RAVI APPLICANT: DAS, KUMUDA C.
 LENGTH: 10
TYPE: PRT
 ORGANISM: Homo sapiens
 TYPE: PRT
 LENGTH: 105
 Local Similarity
 31 WCGPCK 36
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
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 US-10-032-201B-132
 RESULT 55
 US-10-032-201B-123
 US-10-032-201B-123
 FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 132
LENGTH: 106
 Sequence 132, Application US/10032201B Publication No. US20030167524A1
 PILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 123
LENGTH: 106
 Query Match
Best Local Similarity
Query Match
Best Local Similarity
 Query Match
 Publication No. US20030167524A1 GENERAL INFORMATION:
 APPLICANT: Del val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
 APPLICANT:
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
 APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
 APPLICANT:
 TYPE: PRT ORGANISM: Penicillium chrysogenum
 TYPE: PRT
ORGANISM: Helicobacter pylori
 / Match 100.0%; Score 47; DB Local Similarity 100.0%; Pred. No. 12; nes 6; Conservative 0; Mismatches
 31 WCGPCK 36
 29 WCGPCK 34
 1 WCGPCK 6
 1 WCGPCK 6
 6; Conservative
 Deckers, Harm
Heifetz, Peter Bernard
Briggs, Steven
Dalmia, Bipin Kumar
Del Val, Greg
Zaplachinski, Steve
 Briggs, Steven
Dalmia, Bipin Kumar
Del Val, Greg
Zaplachinski, Steve
 Deckers, Harm
Heifetz, Peter Bernard
 Application US/10032201B
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOPTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 176
LENGTH: 106
TYPE: PRT
ROBANISM: Manduca Bexta
US-10-032-201B-176
 Query Match
Best Local Similarity
"---hes 6; Conserv?
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 US-10-032-201B-176
 RESULT 57
 ORGANISM: Streptomyces aureofaciens US-10-032-201B-137
 US-10-032-201B-137
 RESULT 56
 Query Match
Best Local S
 CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 137
LENGTH: 106
 GENERAL INFORMATION:
APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
APPLICANT: Briggs, Steven
 Sequence 137, Application US/10032201B Publication No. US20030167524A1
 Matches
 Sequence 176, Application US/10032201B Publication No. US20030167524A1
 GENERAL INFORMATION
 Matches
 APPLICANT:
APPLICANT:
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
 APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION
TITLE OF INVENTION: COMPOSITIONS
 APPLICANT: MOLONEY, MAUTICE
TITLE OF INVENTION: METHODS FOR THE PRODUCTION
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
 APPLICANT:
 APPLICANT:
 APPLICANT:
 FILE REFERENCE: 38814 351B
 TYPE: PRT
 30 WCGPCK 35
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
 Similarity 6; Conserva
 6; Conservative
 Briggs, Steven
Dalmia, Bipin Kumar
 Deckers, Harm
Heifetz, Peter Bernard
 Briggs, Steven
Dalmia, Bipin Kumar
 Deckers, Harm
Heifetz, Peter Bernard
 Del Val, Greg
Zaplachinski, Steve
 Conservative
 Conservative
 100.0%; Score 47; DB 100.0%; Pred. No. 12;
 100.0%; Score 47;
100.0%; Pred. No.
 0
 0
 0
 Mismatches
 Mismatches
 Mismatches
 4; Length 106
 4; Length 106;
 OF MULTIMERIC
 OF MULTIMERIC
 0
 Indels
 Indels
 Indels
 PROTEINS,
 PROTEINS,
 <u>,</u>
 <u>,</u>
 0,
 Gaps
 Gaps
 Gaps
 AND RELATED
 AND RELATED
 0
 0
 0
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31

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RESULT 59
US-10-628-391-1
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 US-10-335-977-6752
 US-10-335-977-6752
 Sequence 1, Application US/10628391 Publication No. US20040146526A1 GENERAL INFORMATION:
 Sequence 6752, Application US/10335977

Publication No. US20040052799A1

Publication No. US20040052799A1

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: DOUGLAS SMITH et al
APPLICANT: DOUGLAS SMITH et al
APPLICANT: DOUGLAS SMITH et al
APPLICANT: NUCLEIC ACID AMINO ACID SEQUENCES
RELATING TO HELICOBACTER PYLORI FOR
DIAGNOSTICS AND THERAPEUTICS
 Matches
 Query Match
 APPLICANT: WINDLE, Henry J.
APPLICANT: O'TOOLE, Dermot
APPLICANT: KELLEHOR, Dermot
APPLICANT: ABDEL-LATIF, Mohamed M.
TITLE OF INVENTION: INHIBITION OF NF-kappaB ACTIVATION
 TELEPHONE: (617)227-7400
TELEPAX: (617)742-4214
INFORMATION FOR SEQ ID NO: 6752:
SEQUENCE CHARACTERISTICS:
 Local Similarity 100 nes 6; Conservative
 APPLICATION NUMBER: US/10/335,977
FILING DATE: 30-Dec-2002
PRIOR APPLICATION DATA:
 SOPTWARE: UNIX
 APPLICATION NUMBER: 08/993,002
FILING DATE: 17-DEC--997
ATTORNEY/AGENT INFORMATION:
NAME: Mandragouras, Amy E.
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: GTN-018
TELECOMMUNICATION INFORMATION:
 NAME/KEY: misc_feature
LOCATION: (B) LOCATION 1...106
SEQUENCE DESCRIPTION: SEQ ID NO: 6752:
 HYPOTHETICAL: YES
ORIGINAL SOURCE:
ORGANISM: Helicobacter pylori
 COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM 1S09660
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: Windows NT 4.0
 NUMBER OF SEQUENCES: 10031
 FEATURE:
 MOLECULE TYPE:
 CORRESPONDENCE ADDRESS:
 29
 1 WCGPCK 6
 WCGPCK 34
 LENGTH: 106 amino acids TYPE: amino acid TOPOLOGY: linear
 STATE: Massachusetts
 CITY: Boston
 STREET: 28 State Street
 COUNTRY: USA
 ADDRESSEE: LAHIVE & COCKFIELD
 APPLICATION DATA:
 02109-1875
 protein
 100.0%;
 0; Mismatches
 Score 47;
Pred. No.
 ВB
 4.
 0;
 Length 106;
 Indels
 <u>.</u>
 Gaps
 0
```

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APPLICANT: Venter, J. Craig
APPLICANT: Venter, J. Craig
APPLICANT: Vet et al.
TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,
TITLE OF INVENTION: DROSOPHILA GENES.
FILE REFERENCE: CL000728
CURRENT APPLICATION NUMBER: US/11/097,143
CURRENT APPLICATION NUMBER: 00/157,832
PRIOR APPLICATION NUMBER: 60/157,832
PRIOR PILLING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: 60/160,191
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR FILING DATE: 1999-10-28
 US-11-097-143-7479
Sequence 7479, Application US/11097143
Publication No. US20050208558A1
GENERAL INFORMATION:
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 ; OTHER INFORMATION: Clone ID: MRT4577_96874C.1.pep US-10-425-115-365910
 밁
 S
 ; TYPE: PRT; ORGANISM: H. pylori US-10-628-391-1
 RESULT 61
 RESULT 60
 US-10-425-115-365910
 NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 365910
LENGTH: 106
 GENERAL INFORMATION:
APPLICANT: La ROSA, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
 SOFTWARE: PatentIn version 3.2 SEQ ID NO 1
 Query Match
Best Local Similarity
Matches 6; Conserv
 Sequence 365910, Application US/10425115 Publication No. US20040214272A1
 Best Local Similarity
Matches 6; Conserv
 Query Match
 TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants FILE REFERENCE: 38-21(53222)B CURRENT APPLICATION NUMBER: US/10/425,115 CURRENT FILING DATE: 2003-04-28
 CURRENT APPLICATION NUMBER: US/10/628,391
CURRENT FILING DATE: 2003-07-29
PRIOR APPLICATION NUMBER: PCT/IE02/00011
PRIOR FILING DATE: 2002-01-30
NUMBER OF SEQ ID NOS: 9
 FILE REFERENCE: P69048US0
 TYPE: PRT
ORGANISM: Zea mays
FEATURE:
 ENGTH:
 29 WCGPCK 34
 31 WCGPCK 36
 1 WCGPCK 6
 106
 1 WCGPCK 6
 100.0%;
ilarity 100.0%;
Conservative C
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 100.0%;
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 ; Score 47; DB; Pred. No. 12; 0; Mismatches
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 Score 47; DB Pred. No. 12;
 Mismatches
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 Gaps
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RESULT 63
US-10-022-201B-121
; Sequence 121, Application US/10032201B
; Publication No. US20030167524A1
; GENERAL INFORMATION:
; APPLICANT: Von Rooijen, Gijs
; APPLICANT: Deckers, Harm
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 8
 US-10-032-201B-109
 US-10-032-201B-109
 PRIOR APPLICATION NUMBER: 60/173,383
PRIOR FILING DATE: 1999-12-28
PRIOR PELICATION NUMBER: 60/175,693
PRIOR APPLICATION NUMBER: 60/184,831
PRIOR APPLICATION NUMBER: 60/184,831
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: 60/191,637
PRIOR APPLICATION NUMBER: 60/191,637
PRIOR PILING DATE: 2000-03-23
NUMBER OF SEQ ID NOS: 43008
 ; ORGANISM: DROSOPHILA US-11-097-143-7479
 CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 109
LENGTH: 107
 Query Match
Best Local Similarity
Matches 6; Conserv
 GENERAL INFORMATION:
 Sequence 109, Application No.
 Query Match
Best Local S
 SOFTWARE: FastSEQ for Windows Version SEQ ID NO 7479
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter B
 APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: MCTHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
 FILE REFERENCE: 38814 351B
 NAME/KEY: VARIANT
LOCATION: 17, 38, 42, 55, 58, 60, 72, 107
OTHER INFORMATION: Xaa = Any Amino Acid
 LENGTH: 106
TYPE: PRT
 ORGANISM: Chromatium vinosum
 TYPE: PRT
 FEATURE:
 Y MALCH 100.0%;
Local Similarity 100.0%;
les 6; Conservative 0,
 FILING DATE:
 APPLICATION NUMBER: 60/164,769
 31 WCGPCK 36
 31 WCGPCK 36
 1 WCGPCK 6
 WCGPCK 6
 Deckers, Harm
Heifetz, Peter Bernard
 Application US/10032201B o. US20030167524A1
 Conservative
 1999-11-
 100.0%; Score 47; DB
100.0%; Pred. No. 12;
Live 0; Mismatches
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 Score 47; DB 6; Length 106; Pred. No. 12;
 Mismatches
 4; Length 107;
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 Indels
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 Gaps
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 RESULT 64
 Matches
 Matches
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; TYPE: PRT ; ORGANISM: Haemophilus influenzae US-10-032-201B-121
 RESULT 65
US-10-335-977-6753
 ; ORGANISM: Drosophila
US-10-032-201B-151
 US-10-032-201B-151
 Sequence 6753, Application US/10335977
Publication No. US20040052799A1
GENERAL INFORMATION:
APPLICANT: DOUGLAS SMITH et al
TITLE OF INVENTION: NUCLEIC ACII
 APPLICANT: Moloney, Maurice
TITLE OF INVENTION: MOTHOUS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 121
LENGTH: 107
 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 151
 Query Match
Best Local :
 GENERAL INFORMATION:
 Sequence 151, A
Publication No.
 Ouery Match
Best Local
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bernard
APPLICANT: Briggs, Steven
APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
 APPLICANT:
APPLICANT:
 APPLICANT:
 FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
 LENGTH: 107
TYPE: PRT
NUMBER OF SEQUENCES:
 30 WCGPCK 35
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
 Similarity 6; Conserv
 Similarity 6; Conserv
 Briggs, Steven
Dalmia, Bipin Kumar
Del Val, Greg
 Zaplachinski,
 Heifetz,
 Application US/10032201B
o. US20030167524A1
 Conservative
 Conservative
 DOUGLAS SMITH et al
NVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES
NVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES
NVENTION: NUCLEIC AND THERAPEUTICS
 Peter Bernard
 melanogaster
 100.0%;
 100.0%;
 Steve
 10031
 0;
 0,
 Score 47; DB
Pred. No. 12;
0; Mismatches
 Score 47;
Pred. No.
 Mismatches
 DB
12;
 В
 4; Length 107;
 4; Length 107;
 0
 Indels
 Indels
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..
 0
 Gaps
 Gaps
 0,
 0
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CORRESPONDENCE ADDRESS:

COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660

02109-1875

STATE: Massachusetts CITY: Boston

STREET: 28 State Street ADDRESSEE: LAHIVE & COCKFIELD

0

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RESULT 66
US-11-097-143-8226
US-11-097-143-8226
Sequence 8226, Application US/11097143
Publication No. US20050208558A1

Publication No. US20050208558A1

GENERAL INFORMATION:
APPLICANT: Venter, J. Craig
APPLICANT: et al.

TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: DARAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
TITLE OF INVENTION: DROSOPHILA GENES.
FILE REFERENCE: CL000728

CURRENT PILING DATE: 1005-04-04
PRIOR APPLICATION NUMBER: 60/157,832
PRIOR APPLICATION NUMBER: 60/157,832
PRIOR APPLICATION NUMBER: 60/160,191
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR FILING DATE: 1999-10-19
PRIOR FILING DATE: 1999-10-19
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR PILING DATE: 1999-10-19
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR PILING DATE: 1999-10-28
PRIOR APPLICATION NUMBER: 60/161,769
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 SEQUENCE DESCRIPTION: SEQ ID NO: 6753: US-10-335-977-6753
 Query Match
Best Local Similarity lou.
Conservative
 TELEPHONE: (617)227-7400
TELEPAX: (617)742-4214
INFORMATION FOR SEQ ID NO: 6753:
SEQUENCE CHARACTERISTICS:
 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/335,977
PILING DATE: 30-Dec-2002
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/993,002
PILING DATE: 17-DEC-1997
ATTORNEY/AGENT INFORMATION:
NAME: Mandragouras, Amy E.
REGISTRATION NUMBER: 36,207
FILING DATE: 1999-11-12
APPLICATION NUMBER: 60/173,383
FILING DATE: 1999-12-28
 HYPOTHETICAL: YES ORIGINAL SOURCE:
 REFERENCE/DOCKET NUMBER: GTN-018 TELECOMMUNICATION INFORMATION:
 FEATURE:
 MOLECULE TYPE: protein
 30 WCGPCK 35
 1 WCGPCK 6
 LENGTH: 107 amino acids TYPE: amino acid TOPOLOGY: linear
 NAME/KEY: misc_feature
LOCATION: (B) LOCATION 1...107
 ORGANISM: Helicobacter pylori
 SOFTWARE: UNIX
 COMPUTER: IBM PC Compatible OPERATING SYSTEM: Windows NT 4.0
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 Score 47; DB Pred. No. 12;
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 Indels
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 Gaps
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; PRIOR FILING DATE: 2000-01-12; PRIOR APPLICATION NUMBER: 60/184,831; PRIOR FILING DATE: 2000-02-24; PRIOR APPLICATION NUMBER: 60/191,637; PRIOR FILING DATE: 2000-03-23; NUMBER OF SEQ ID NOS: 43008; SOFTWARE: FastSEQ for Windows Version 4.15 SEQ ID NO 8226; SEQ ID NO 8226
 Sequence 117, Application US/10
Publication No. US20030167524A1
GENERAL INFORMATION:
APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Ber
 RESULT 68
US-10-032-2018-117
 문
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 ; TYPE: PRT; ORGANISM: Escherichia coli US-09-812-777-16
 RESULT 67
US-09-812-777-16
 밁
 Ś
 ; ORGANISM: DROSOPHILA US-11-097-143-8226
 CURRENT APPLICATION NUMBER: US/09/812,777
CURRENT FILING DATE: 2001-03-18
PRIOR APPLICATION NUMBER: KR 00-63596
PRIOR FILING DATE: 2000-10-27
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 16
LENGTH: 108
PILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 117
 Sequence 16, Application US/09812777
Patent No. US20020115827A1
GENERAL INFORMATION:
 Query Match
Best Local (
 Matches
 Query Match 100.0%; Score 47; DB Best Local Similarity 100.0%; Pred. No. 12;
 APPLICANT:
 Matches
 APPLICANT: KIM, KEY-SUN
APPLICANT: LEB, DUCK YEON
TITLE OF INVENTION: A METHOD OF PROTEIN STABILIZATION
PILE REFERENCE: 05823.0195-00000
 APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
 APPLICANT:
 TYPE: PRT
 y Match 100.0%; Score 47; DB 6; Length 107;
Local Similarity 100.0%; Pred. No. 12;
hes 6; Conservative 0; Mismatches 0; Indels
 30 WCGPCK 35
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
 6,
 Briggs, Steven
Dalmia, Bipin Kumar
 Heifetz, Peter Bernard
 Del Val, Greg
Zaplachinski, Steve
 Application US/10032201B
 Conservative
 US20030167524A1
 0; Mismatches

 Length 108

 Indels
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 ٥,
 Gaps
 Gaps
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FILE REFERENCE: 38814 3518
; CURRENT APPLICATION NUMBER: US/10/032,2018
; CURRENT FILING DATE: 2001-12-19
; NUMBER OF SEQ ID NOS: 313
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 133
; LENGTH: 108
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-10-032-2018-133
 밁
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 ; TYPE: PRT ; ORGANISM: E coli, salmonella typhimurium US-10-032-201B-117
 US-10-032-201B-133
 CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 142
LENGTH: 108
 Query Match 100.0%; Score 47; DB 4; Length 108; Best Local Similarity 100.0%; Pred. No. 12; Matches 6; Conservative 0; Mismatches 0; Indels
 Query Match 100.0%; Score 47; DB 4; Length 108; Best Local Similarity 100.0%; Pred. No. 12; Matches 6; Conservative 0; Mismatches 0; Indels
 Sequence 142, Application US/10032201B Publication No. US20030167524A1
 Sequence 133, Application US/10032201B Publication No. US20030167524A1
 -10-032-201B-142
 APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: MCTHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
 APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
 FILE REFERENCE: 38814 3518
 LENGTH: 108
ORGANISM: Thiobacillus ferrooxidans
 INFORMATION:
 31 WCGPCK 36
 32 WCGPCK 37
 1 WCGPCK 6
 Deckers, Harm
Heifetz, Peter Bernard
 Deckers, Harm
Heifetz, Peter Bernard
 <u>.</u>
 0;
 Gaps
 Gaps
 0;
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US-10-032-201B-142

Query Match
Best Local Similarity 100.0%; Score 47; DB 4; Length 108;
Best Local Similarity 100.0%; Pred. No. 12;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps

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Db 32 WCGPCK 37
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Search completed: February 23, 2006, 00:47:55 Job time : 125.5 secs

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Perfect score:
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4: /cgn2-6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

5: /cgn2-6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

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7: /cgn2-6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

8: /cgn2-6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*
 length: 0
length: 2000000000
 US-10-660-118A-3
47
1 WCGPCK 6
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Query
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US-11-131-744-2
US-11-131-744-1
US-11-094-176-286
US-11-090-916-7
US-11-090-916-8
US-11-090-916-10
US-11-090-916-11
US-11-090-916-12
US-11-090-916-12
US-11-106-796-13
US-11-104-796-13
US-11-090-916-4
US-11-090-916-4
US-11-106-796-12
US-11-090-916-4
US-11-106-796-12
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APPLICANT: White, Carl W.
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OTHER INFORMATION: synthetic peptide motif
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TITLE OF INVENTION: SPECTROMETRY

FILE REFERENCE: NVX-28 US

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APPLICANT: SADEGHI, MEHRNOOSH
APPLICANT: LIANG, XIQUAN
TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR CALIBRATION IN MASS
TITLE OF INVENTION: SPECTROMETRY
FILE REFERENCE: NVX-28 US
CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT APPLICATION NUMBER: 60/572,290
PRIOR APPLICATION NUMBER: 60/572,290
PRIOR APPLICATION NUMBER: 60/659,373
PRIOR APPLICATION NUMBER: 60/659,373
PRIOR APPLICATION NUMBER: 60/6572,215
PRIOR APPLICATION NUMBER: 60/6572,215
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR FILING DATE: 2004-05-05-17
PRIOR APPLICATION NUMBER: 60/621,686
PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 9
SOFTWARE: PatentIn Ver. 3.3
SEQ ID NO 1
LENGTH: 91
 á
 RESULT 3
US-11-131-744-1
 문
 US-11-131-744-2
 US-11-131-744-2
 Sequence 1, Application US/11131744
Publication No. US20060023808A1
GENERAL INFORMATION:
 Matches
 Query Match
Best Local Similarity
 OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: protein sequence
 FEATURE:
 32 WCGPCK 37
 1 WCGPCK 6
 ς.
 Conservative
 100.0%; Score 47; Di
100.0%; Pred. No. 0.:
cive 0; Mismatches
 0;
 0.2;
 DB 7; Length 75;
 0
 0,
 Gaps
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0,

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; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: protein sequence US-11-131-744-1
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 8
 ; SEQ ID NO 286; LENGTH: 103; TYPE: PRT; ORGANISM: Lactobacillus acidophilus US-11-074-176-286
 밁
 US-11-090-916-7
 US-11-074-176-286
 Sequence 7, Application US/11090916 Publication No. US20050260140A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 Sequence 286, Application US/11074176 Publication No. US20050250135A1
 Matches
 Query Match
Best Local Similarity
 APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
 PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
 PRIOR APPLICATION NUMBER: 60/551,161
PRIOR FILING DATE: 2004-03-08
NUMBER OF SEQ ID NOS: 381
 FILE REFERENCE: 5051-694
CURRENT APPLICATION NUMBER: US/11/074,176
CURRENT FILING DATE: 2005-03-07
 APPLICANT: Peril, Andrea Azcarate
TITLE OF INVENTION: Nucleic Acid Sequences Encoding
TITLE OF INVENTION: Stress-Related Proteins and Uses Therefore
 APPLICANT: Klaenhammer, Todd R. APPLICANT: Russell, William M.
 SOFTWARE: FastSEQ for Windows Version 4.0
 APPLICANT:
 TYPE: PRT ORGANISM: Artificial Sequence
 FEATURE:
 FILING DATE: 2004-03-24
APPLICATION NUMBER: 60/650,865
 27 WCGPCK 32
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
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 9
 Altermann, Eric
McAuliffe, Olivia
 Conservative
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2005-02-07
 100.0%; Score 47; DB 7; Length 103; 100.0%; Pred. No. 0.25;
 100.0%;
 0
 Score 47; DB 7; Length 91
Pred. No. 0.23;
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 Indels
 0;
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 Gaps
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 Query Match
Best Local Similarity
""" hes 6; Conserve
 ; LENGTH: 103
TYPE: PRT
; ORGANISM: Saccharomyces cerevisiae
US-11-090-916-7
 US-11-090-916-8
 ; ORGANISM: Homo sapiens US-10-821-234-1371
 RESULT 6
 US-10-821-234-1371
 Sequence 8, Application US/11090916 Publication No. US20050260140A1 GENERAL INFORMATION:
 NUMBER OF SEQ ID NOS: 1704
SOPTWARE: pt_SEQ_genes Version 1.0
SEQ ID NO 1371
 SOFTWARE: PatentIn version 3.3 SEQ ID NO 7
 Matches
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 GENERAL INFORMATION:
 Sequence 1371, Application US/10821234
Publication No. US20050255114A1
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR PRIOR DATE: 2003-09-11
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/560,118
PRIOR APPLICATION NUMBER: 10/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR PILING DATE: 2003-02-07
NUMBER OF SEO ID NOS: 15
 FILE REFERENCE: 821A
CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
 APPLICANT: Labat, Ivan APPLICANT: Stache-Cra
 APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
 APPLICANT:
 TYPE: PRT
 ENGTH: 105
 Local Similarity
 31 WCGPCK 36
 29 WCGPCK 34
 1 WCGPCK 6
 1 WCGPCK 6
 6,
 Stache-Crain, Birgit
 del Val, Greg
 Andarmani, Susan
 100.0%; Score 47; DB 7 illarity 100.0%; Pred. No. 0.25; Conservative 0; Mismatches
 Conservative
 Carl W.
 100.0%;
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 Score 47; DB 6
Pred. No. 0.25;
 Mismatches
 DB 6; Length 105;
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 Indels
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밁
 ; TYPE: PRT ; ORGANISM: Mus musculus US-11-090-916-9
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 ; ORGANISM: Gallus gallus US-11-090-916-8
 US-11-090-916-10
 RESULT 9
 US-11-090-916-9
 PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR PILING DATE: 2004-03-24
 Sequence 10, Application US/11090916 Publication No. US20050260140A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.3
SEQ ID NO 9
LENGTH: 105
 Sequence 9, Application US/11090916 Publication No. US20050260140A1
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 Query Match
Best Local Similarity
 GENERAL INFORMATION:
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 Query Match
 TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
FRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR FILING DATE: 2002-09-10
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 60/460,118
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/560,516
PRIOR APPLICATION NUMBER: 10/556,516
 APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: del, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
 APPLICANT: White, Carl W. APPLICANT: del Val, Greg APPLICANT: Lee, Rees
 NUMBER OF SEQ ID NOS: 15
 PRIOR APPLICATION NUMBER: 60/650,865 PRIOR FILING DATE: 2005-02-07
 TYPE:
 LENGTH:
 Local Similarity les 6; Conserv
FILING DATE:
 31 WCGPCK 36
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
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 Lee, Rees
 100.0%; ilarity 100.0%; Conservative 0
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 Carl W.
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 Score 47; DB 7
Pred. No. 0.25;
 Score 47; DB 7
Pred. No. 0.25;
 Mismatches
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 DB 7; Length 105;
 DB 7; Length 105;
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 Indels
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 Gaps
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; TYPE: PRT; ORGANISM: Bos taurus US-11-090-916-11
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 RESULT 10
US-11-090-916-11
 US-11-090-916-12
 US-11-090-916-10
 PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILLING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
PRIOR FILING DATE: 2004-03-24
 NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin version 3.3
SEQ ID NO 10
LENGTH: 105
 Sequence 12, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
 SEQ ID NO 11
 GENERAL INFORMATION:
 Sequence 11, Application US/11090916
Publication No. US20050260140A1
 Matches
 Best Local Similarity
 Query Match
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 Query Match
 APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
 APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
 PRIOR APPLICATION NUMBER: 60/650,865 PRIOR FILING DATE: 2005-02-07
 NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn versi
 PRIOR FILING DATE:
 APPLICANT: White,
 PRIOR APPLICATION NUMBER: 60/650,865
PRIOR APPLICATION NUMBER: 60/462,082
 ORGANISM: Rattus norvegicus
 TYPE: PRT
 LENGTH:
 Local
 31 WCGPCK 36
 31 WCGPCK
 1 Similarity 6; Conserv
 1 WCGPCK 6
 105
 1 WCGPCK 6
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 PatentIn version 3.3
 Conservative
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 Carl W.
 2005-02-07
 100.0%;
 100.0%; Score 47;
100.0%; Pred. No.
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 Score 47;
Pred. No.
 Mismatches
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 0.25;
 DB 7; Length 105;
 DB 7;
 Length 105;
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 Gaps
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PRIOR FILING DATE: 2003-04-11

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RESULT 13
US-11-144-301A-9
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 RESULT 12
US-11-106-796-13
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 US-11-106-796-13
 US-11-090-916-12
 Sequence 13, Application US/11106796
Publication No. US20050282191A1
GENERAL INFORMATION:
APPLICANT: SUTOYSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
FILE REFERENCE: UVMO:033US
CURRENT EPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2005-04-15
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
SOCTWARE: Patentin Ver. 2.1
SEQ ID NO 13
 Sequence 9, Application US/11144301A Publication No. US20050288227A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
Matches 6; Conserv
 Best Local Similarity
Matches 6; Conserv
 SOFTWARE: PatentIn version 3.3 SEQ ID NO 12
 Query Match
 APPLICANT: Johanna, Ungerstedt
TITLE OF INVENTION: USE OF THLOREDOXIN MEASUREMENTS FOR DIAGNOSTICS AND TREATMENTS
FILE REPERENCE: 28617-501 UTIL
CURRENT APPLICATION NUMBER: US/11/144,301A
CURRENT FILING DATE: 2005-06-03
PRIOR APPLICATION NUMBER: US/577,089
PRIOR APPLICATION NUMBER: US 60/577,089
PRIOR FILING DATE: 2004-06-04
PRIOR FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR APPLICATION NUMBER: US 60/357,383
PRIOR FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 10
 APPLICANT: Marks, Paul A. APPLICANT: Johanna, Unge
 WHIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
 PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
 PRIOR APPLICATION NUMBER: 10/660,118 PRIOR FILING DATE: 2003-09-10
 LENGTH: 105
TYPE: PRT
ORGANISM: Homo sapiens
 TYPE: PRT
ORGANISM: Homo sapiens
 ENGTH: 105
 FILING DATE: 2003-09-10
APPLICATION NUMBER: 60/556,516
FILING DATE: 2004-03-24
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 31
 1 WCGPCK 6
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 WCGPCK 36
 WCGPCK 6
 WCGPCK 36
 100.0%; Score 47; DB 7 ilarity 100.0%; Pred. No. 0.25; Conservative 0; Mismatches
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 Score 47; DB 7
Pred. No. 0.25;
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 DB 7;
 DB 7; Length 105;
 0,
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 Length 105
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 0,
 Gaps
 Gaps
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APPLICANT: Levy, Ronald
TITLE OF INVENTION: DESIGN OF THERAPEUTICS AND THERAPEUTICS
FILE REFERENCE: 17102-013001 / 1762
FULE REFERENCE: 17102-013001 / 1762
CURRENT APPLICATION NUMBER: US/11/032,773
CURRENT FILING DATE: 2005-01-11
PRIOR APPLICATION NUMBER: 60/536,184
PRIOR FILING DATE: 2004-01-12
PRIOR FILING DATE: 2004-03-29
1 NUMBER OF SEQ ID NOS: 958
1 SOFTWARE: FastSEQ for Windows Version 4.0
1 SEQ ID NO 943
1 LENGTH: 105
 RESULT 15
US-11-032-773-943
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 Query Match
Best Local Similarity
Thes 6; Conserve
 US-11-144-301A-9
 US-11-144-301A-10
 CURRENT APPLICATION NUMBER: US/11/144,301A
CURRENT FILING DATE: 2005-06-03
PRIOR APPLICATION NUMBER: US 60/577,089
PRIOR FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR FILING DATE: 2003-02-14
PRIOR PRIOR PRIOR DATE: 2003-02-15
PRIOR FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 10
 US-11-144-301A-10
 Sequence 943, Application US/11032773 Publication No. US20060018911A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.3 SEQ ID NO 10
 SEQ ID NO 9
 Matches
 Query Match
 GENERAL INFORMATION:
 Sequence 10, Application US/11144301A Publication No. US20050288227A1
 TYPE: PRT
ORGANISM: Mus musculus
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank BAA04881
 APPLICANT: Marks, Paul A. APPLICANT: Johanna, Unge
 FILE REFERENCE: 28517-501 UTIL
 APPLICANT: JOHANNA, UNGERSTEELT MEASUREMENTS FOR DIAGNOSTICS AND TREATMENTS
 LENGTH: 105
TYPE: PRT
ORGANISM: Homo sapiens
 TYPE: PRT
ORGANISM: Homo sapiens
DATABASE ENTRY DATE: 2002-12-25
 ENGTH:
 y Match 100.0%;
Local Similarity 100.0%;
les 6; Conservative 0
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 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
 105
 Conservative
 100.0%;
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 Score 47; DB 7
Pred. No. 0.25;
 Score 47; DB 7; Pred. No. 0.25;); Mismatches (
 DB 7; Length 105;
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 Length 105
 Indels
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 ; TYPE: PRT; ORGANISM: Pseudomonas syringae US-11-090-916-4
 US-11-106-796-11
 FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR PPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/56,516
PRIOR FILING DATE: 2003-09-10
PRIOR PPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR APPLICATION NUMBER: 60/650,865
 US-11-090-916-4
 RESULT 16
 US-11-032-773-943
 SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 11
LENGTH: 127
 PRIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.:
SEQ ID NO 4
 Query Match
Best Local Similarity
 Sequence 11, Application US/11106796
Publication No. US20050282191A1
GENERAL INFORMATION:
 Matches
Query Match
Best Local Similarity
 Matches
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 Sequence 4, Application US/11090916
Publication No. US20050260140A1
 FILE REFERENCE: UVMO:033US
CURRENT APPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2005-04-15
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
 APPLICANT: White, Carl W.
APPLICANT: del Val. Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
 TYPE: PRT
ORGANISM: Mus musculus
.11-106-796-11
 APPLICANT: SUTOVSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
 LENGTH: 109
 31 WCGPCK 36
 33 WCGPCK 38
 1 WCGPCK 6
 1 WCGPCK 6
 6;
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 Conservative
 Conservative
 version 3.3
 100.0%; Score 47; DB 7
100.0%; Pred. No. 0.26;
100.0%;
 100.0%; Score 47; I
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Score 47;
Pred. No.
 Mismatches
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 DB 7;
 DB 7; Length 105;
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 Length 109
 Length 127;
 Indels
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 Gaps
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RESULT 20
US-11-131-744-3
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 Query Match
Best Local Similarity
Thebe 6; Conserv
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 ; ORGANISM: RAT
US-11-106-796-12
 ; ORGANISM: Neisseria gonorrhoeae US-10-467-657-4354
 US-11-106-796-12
 US-10-467-657-4354
 SOPTWARE: SeqWin99, version 1.04
SEQ ID NO 4354
LENGTH: 129
 Sequence 4354, Application US/10467657 Publication No. US20050260581A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn
SEQ ID NO 12
LENGTH: 127
 Sequence 12, Application US/11106796 Publication No. US20050282191A1 GENERAL INFORMATION:
 Matches
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 Query Match
 Best Local Similarity
 APPLICANT: SUTOVSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
FILE REFERENCE: UVMO:033US
 FILE REFERENCE:

CURRENT APPLICATION NUMBER: US/10/467,657

CURRENT FILING DATE: 2003-08-11

PRIOR APPLICATION NUMBER: GB-0103424.8

PRIOR FILING DATE: 2001-02-12

NUMBER OF SEQ ID NOS: 9218
 APPLICANT: CHIRON SPA
APPLICANT: FONTANA Maria Rita
APPLICANT: PIZZA Mariagrazia
APPLICANT: MASIGNANI Vega
APPLICANT: MONACI Elisabetta
 PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
 CURRENT APPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2005-04-15
 TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS
 LENGTH: 12
TYPE: PRT
 TYPE: PRT
 31
 31 WCGPCK 36
 52 WCGPCK 57
 1 WCGPCK 6
 1 WCGPCK 6
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 100.0%; Score 47; DB 7; Length 127; ilarity 100.0%; Pred. No. 0.29; Conservative 0; Mismatches 0; Indels
 Ver. 2.1
 100.0%; Score 47; DB 6; 100.0%; Pred. No. 0.29;
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 Mismatches
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 Indels
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 Gaps
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; Sequence 3, Application US/11131744

Publication No. US20060023808A1

APPLICANT: HAJIVANDI, MAHBOD R

SADEGHI, MEHRNOOSH

ROBERT M.

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밁
 US-11-131-744-6
 US-11-131-744-3
 SOFTWARE: PatentIn
SEQ ID NO 6
 Sequence 6, Application US/11131744 Publication No. US20060023808A1
 Best Local Similarity
Matches 6; Conserv
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 SEQ ID NO 3
 GENERAL INFORMATION
 PRIOR APPLICATION NUMBER: 60/572,290 PRIOR FILING DATE: 2005-05-17 PRIOR APPLICATION NUMBER: 60/669,373 PRIOR FILING DATE: 2005-04-08 PRIOR APPLICATION NUMBER: 60/572,215 PRIOR FILING DATE: 2004-05-17
 PRIOR APPLICATION NUMBER: 60/669,373
PRIOR FILING DATE: 2005-04-08
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR FILING DATE: 2004-05-17
 PRIOR APPLICATION NUMBER: 60/572,290 PRIOR FILING DATE: 2005-05-17
 CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
 TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR CALIBRATION IN MASS TITLE OF INVENTION: SPECTROMETRY FILE REFERENCE: NVX-28 US
 APPLICANT:
 APPLICANT: HAJIVANDI, MAHBOD R. APPLICANT: POPE, ROBERT M.
 NUMBER OF SEQ ID NOS: 9
 CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
 APPLICANT: LEITE, JOHN F.
TITLE OF INVENTION: COMPOSITIONS,
TITLE OF INVENTION: SPECTROMETRY
 PRIOR FILING DATE: 2004-10-26
 APPLICANT:
 SOFTWARE: PatentIn Ver. 3.3
 NUMBER OF SEQ
 FILE REFERENCE: NVX-28 US
 FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: protein sequence
 ORGANISM: Artificial Sequence
DRGANISM: Artificial Sequence
 ENGTH:
 ENGTH:
 FILING DATE: 2004-05-17
APPLICATION NUMBER: 60/621,685
 APPLICATION NUMBER: 60/621,686
 APPLICATION NUMBER: 60/621,685
FILING DATE: 2004-10-26
 FILING DATE: 2004-10-26
 APPLICATION NUMBER: 60/621,686
 31 WCGPCK 36
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 1 WCGPCK 6
 SADEGHI, MEHRNOOSH
 ID NOS:
 Conservative
 Ver. 3.3
 100.0%; Score 47;
100.0%; Pred. No.
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 KITS, AND METHODS FOR CALIBRATION IN MASS
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 Indels
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 Gaps
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RESULT 23
US-11-090-916-6
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 ; PEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic;
; OTHER INFORMATION: protein sequence
US-11-131-744-6
Sequence 6, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rese
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
 Query Match
Best Local Similarity
 SOFTWARE: PatentIn Ver. 3.3 SEQ ID NO 7
 Sequence 7, Application US/11131744
Publication No. US20060023808A1
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 Best Local Similarity Matches 6; Conserv
 Query Match
 GENERAL INFORMATION
 APPLICANT:
 CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/572,290
PRIOR FILING DATE: 2005-05-17
 PRIOR APPLICATION NUMBER: 60/621,686 PRIOR FILING DATE: 2004-10-26 NUMBER OF SEQ ID NOS: 9
 APPLICANT: HAJIVANDI, MAHBOD R. APPLICANT: POPE, ROBERT M.
 PRIOR APPLICATION NUMBER: 60/669,373
PRIOR FILING DATE: 2005-04-08
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR FILING DATE: 2004-05-17
 TITLE OF INVENTION: COMPOSITIONS, TITLE OF INVENTION: SPECTROMETRY FILE REFERENCE: NVX-28 US
 TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
 OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: protein sequence
 APPLICANT:
 LENGTH: 440
 APPLICATION NUMBER: 60/621,685 FILING DATE: 2004-10-26
 32 WCGPCK 37
 31
 1 WCGPCK 6
 1 WCGPCK 6
 6
 LIANG,
 LEITE, JOHN F.
 SADEGHI, MEHRNOOSH
 Conservative
 Conservative
 100.0%;
 100.0%; Score 47; DB 7
100.0%; Pred. No. 0.65;
 0
 Score 47;
Pred. No.
 Mismatches
 KITS, AND METHODS FOR CALIBRATION IN MASS
 DB 7; Length 440
 DB 7;
 <u>,</u>
 Length 273
 <u>,</u>
 0
 Gaps
 Gaps
 0
 0
```

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RESULT 24
US-11-090-916-5
 S
 Query Match
Best Local Similarity
Watches 5; Conserv
 ; LENGTH: 103
TYPE: PRT
; ORGANISM: Listeria monocytogenes
US-11-090-916-6
 RESULT 25
US-11-098-686-10612
 밁
 8
 US-11-090-916-5
 PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR PELLING DATE: 2003-09-10
PRIOR PELLING DATE: 2003-04-11
PRIOR PELLING DATE: 2003-04-11
PRIOR PELLING DATE: 2003-09-10
PRIOR PELLING DATE: 2003-09-10
PRIOR PELLING DATE: 2004-03-24
PRIOR PELLING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR PELLING DATE: 2005-02-07
 Sequence 10612, Application US/11098686
publication No. US20060024696A1
GENERAL INFORMATION:
APPLICANT: Kapur, Vivek and Gebhart, Connie J.
TITLE OF INVENTION: NUCLEIC ACID AND POLYPEPTIDE SEQUENCES
TITLE OF INVENTION: FROM LAWSONIA INTRACELLULARIS AND METHODS OF USING
 NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin version 3.3
SEQ ID NO 5
LENGTH: 104
 Sequence 5, Application US/11090916 Publication No. US20050260140A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
Matches 5; Conserv
 SOFTWARE: PatentIn version 3.3 SEQ ID NO 6
 APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: del Res
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
CURRENT FILING DATE: 2005-03-24
 PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
 PRIOR APPLICATION NUMBER: 60/556,516 PRIOR FILING DATE: 2004-03-24
 PRIOR PRIOR
 NUMBER OF SEQ ID NOS: 15
 PRIOR APPLICATION NUMBER: 60/462,082 PRIOR FILING DATE: 2003-04-11
FILE REFERENCE:
 TYPE: PRT ORGANISM: Porphyromonas gingivalis
 APPLICATION NUMBER: 10/660,118 FILING DATE: 2003-09-10
 27 WCGPCR 32
 28 WCGPCR 33
 1 WCGPCK 6
 1 WCGPCK 6
 Conservative
 Conservative
 93.6%;
83.3%;
 93.6%;

 Mismatches

 Score 44; DB 7;
Pred. No. 0.74;
 Score 44; DB 7
Pred. No. 0.74;
 Mismatches
 DB 7;
 0
 0; Indels
 Length 104;
 Length 103;
 0
 0
 Gaps
 Gaps
 0
 0;
```

```
Sequence 12, Application US/11195459
Publication No. US20050278803A1
GENERAL INFORMATION:
APPLICANT: Sewalt, Vincent
APPLICANT: Hastings, Craig
APPLICANT: Hestings, Craig
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
APPLICANT: Jung, Rudolf
 ; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 10612
; LENGTH: 107
; TYPE: PRT
; ORGANISM: Lawsonia intracellularis
US-11-098-686-10612
 밁
 á
 US-11-195-459-16
Sequence 16, Application US/11195459
Publication No. US20050278803A1
GENERAL INFORMATION:
 밁
 Ś
 RESULT 26
US-11-195-459-12
 RESULT 27
 US-11-195-459-12
 Best Loc
Matches
 NUMBER OF SEQ ID NOS 25
SOFTWARE: PatentIn version 3.0
SEQ ID NO 12
LENGTH: 122
 Query Match
 Query Match
Best Local Similarity
 Matches
 APPLICANT:
 APPLICANT:
APPLICANT:
 CURRENT APPLICATION NUMBER: US/11/098,686
CURRENT FILLING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US03/31318
PRIOR FILING DATE: 2003-10-01
PRIOR APPLICATION NUMBER: US 60/416,395
PRIOR FILING DATE: 2002-10-04
NUMBER OF SEQ ID NOS: 11433
 APPLICANT: Jung, Rudolf
APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PRC
FILE REFERENCE: 5718-119 (035718/241421)
CURRENT PEPLICATION NUMBER: US/11/195, 459
CURRENT FILING DATE: 2005-08-02
 APPLICANT: Sewalt, Vincent APPLICANT: Hastings, Crai
 PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
 APPLICANT:
 TYPE: PRT
ORGANISM: Zea mays
 Local
 39 WCGPCR 44
 29 WCGPCR
 1 WCGPCK 6

 7: Jung, Rudolf
 F. Everard, John
 F. Allen, Stephen
 INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS

 1 WCGPCK 6
 Similarity 5; Conserv
 Hastings, Craig
Meeley, Robert
Hantke, Sabine
Jung, Rudolf
 Conservative
 Conservative
 34
 Craig
 93.6%;
 93.6%; Score 44; DB 7;
83.3%; Pred. No. 0.76;
rive 1; Mismatches
 Score 44; DB 7
Pred. No. 0.82;
 Mismatches
 DB 7;
 0
 0;
 Length 122;
 Length 107;
 Indels
 Indels
 <u>,</u>
 0
 Gaps
 Gaps
 0
 0,
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PR

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CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin version 3.0
SEQ ID NO 16
LENGTH: 122
TYPE: PRT
ORGANISM: Hordeum vulgare
 RESULT 29
US-11-195-459-10
 ; FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: "n" at position 9,
OTHER INFORMATION: or t
US-11-195-459-19
 APPLICANT: Hastings, Craig
APPLICANT: Meeley, Robert
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
APPLICANT: Jung, Rudolf
APPLICANT: Sverard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
FILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR APPLICATION NUMBER: 00/250,703
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
 S
 문
 RESULT 28
US-11-195-459-19
 밁
 US-11-195-459-16
 Sequence 10, Application US/11195459 Publication No. US20050278803A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 19 LENGTH: 122
 Matches
 Sequence 19, Application US/11195459 Publication No. US20050278803A1
 Query Match
Best Local Similarity
 Query Match
Best Local
 GENERAL INFORMATION:
 Matches
 APPLICANT: Sewalt, Vincent APPLICANT: Hastings, Craig
 APPLICANT: Sewalt, Vincent
 FILE REFERENCE: 5718-119 (035718/241421)
 APPLICANT:
 ORGANISM: Zea mays
 TYPE: PRT
 39 WCGPCR 44
 45
 1 WCGPCK 6
 1 WCGPCK 6
 ٠.
ت
 Similarity
 WCGPCR 50
Hastings, Crai
Meeley, Robert
 Conservative
 Conservative
 93.6%;
 93.6%;

 Mismatches

 Score 44; DB 7;
Pred. No. 0.82;
1; Mismatches
 Score 44; DB 7
Pred. No. 0.82;
 493,
 DB 7;
 537,
 FOR ALTERING THE DISULFIDE STATUS OF
 <u>..</u>
 Length 122
 548,
 Length 122
 Indels
 581,
 and 584 can be
 0
 0,
 Gaps
 Gaps
 0
 an
 0;
 a,
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APPLICANT: White, Carl W.
APPLICANT: Hel Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/56,516
PRIOR APPLICATION NUMBER: 60/560,865
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
RESULT 31
US-11-195-459-8
 문
 Query Match
Best Local Similarity
Thehes 5; Conservi
 밁
 ; ORGANISM: Zea mays
US-11-195-459-10
 ; ORGANISM: Arabidopsis thaliana US-11-090-916-13
 US-11-090-916-13
 Query Match
Best Local S
Matches 5
 SOFTWARE: PatentIn version 3.3
SEQ ID NO 13
LENCTH: 134
TYPE: PRT
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 10
 GENERAL INFORMATION:
 Sequence 13, Ap
Publication No.
 APPLICANT: Jung, Rudolf
APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS
FILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR PILING DATE: 2001-12-03
PRIOR APPLICATION NUMBER: 60/250,703
PRIOR APPLICATION STATE CONTINUED NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
 NUMBER OF SEQ ID NOS: 15
 LENGTH: 123
TYPE: PRT
 Local Similarity
 40 WCGPCR 45
 59 WCGPCR
 1 WCGPCK 6
 1 WCGPCK 6
 Application US/11090916
No. US20050260140A1
 Hantke, Sabine
 Conservative
 Conservative
 64
 93.6%;
 93.6%;
 Score 44; DB 7;
Pred. No. 0.88;
 Score 44; I
Pred. No. 0
 Mismatches
 Mismatches
 DB 7;
 0
 0
 Length 123
 Length 134;
 Indels
 0,
 0
 Gaps
 Gaps
 0
 0
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 PR
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Sequence 8, Application US/11195459
Publication No. US20050278803A1

GENERAL INFORMATION:

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RESULT 33
US-11-090-916-15
 RESULT 32
US-11-090-916-14
 Query Match
Best Local Similarity
Tatches 5; Conserv
 CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/560,516
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR PILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR APPLICATION NUMBER: 60/650,865
 밁
 ; TYPE: PRT; ORGANISM: Zea mays US-11-195-459-8
 밁
 Ś
 US-11-090-916-14
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 8
 Query Match
Best Local Similarity
Matches 5; Conserv
 NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.3
SEQ ID NO 14
LENGTH: 167
 Sequence 14, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
 APPLICANT: Hastings, Craig
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
APPLICANT: Hantke, Sabine
APPLICANT: Jung, Rudolf
APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PROFILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT APPLICATION NUMBER: US/10/005,429
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR APPLICATION NUMBER: 60/250,703
 PRIOR FILING DATE: 2000-: NUMBER OF SEQ ID NOS: 25
 TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof FILE REFERENCE: 2879-98-1
 APPLICANT: White, Carl W. APPLICANT: del Val, Greg
 PRIOR FILING DATE:
 APPLICANT: Lee, Rees
 APPLICANT: Sewalt, Vincent
 TYPE: PRT
ORGANISM: Zea mays
 LENGTH: 134
 54 WCGPCR 59
 88 WCGPCR 93
 1 WCGPCK 6
 1 WCGPCK 6
 Conservative
 Conservative
 Carl W.
 2000-12-01
 2005-02-07
 93.6%;
 93.6%;

 Mismatches

 Score 44; D
Pred. No. 1;
 Score 44; DB 7;
Pred. No. 0.88;
 Mismatches
 0
 0;
 Length 134;
 Indels
 Indels
 0
 <u>,</u>
 Gaps
 Gaps
 0
 ٥,
```

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Sequence 8494, Application US/10467657
Publication No. US2005280581A1
GENERAL INFORMATION:
APPLICANT: CHIRON SpA
APPLICANT: FONTANA Maria Rita
APPLICANT: MASIGNANI Vega
APPLICANT: MASIGNANI Vega
APPLICANT: MONACI Elisabetta
 밁
 á
 ; ORGANISM: Oryza sativa
US-11-090-916-15
 문
 FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILLING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR PILLING DATE: 2003-04-11
PRIOR PILLING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR PILLING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILLING DATE: 2004-03-24
PRIOR FILLING DATE: 2004-03-24
PRIOR FILLING DATE: 2004-03-24
PRIOR FILLING DATE: 2004-03-24
 ; ORGANISM: Neisseria gonorrhoeae US-10-467-657-8494
 US-10-467-657-8494
 CURRENT APPLICATION NUMBER: US/10/467,657
CURRENT FILING DATE: 2003-08-11
PRIOR APPLICATION NUMBER: GB-0103424.8
PRIOR FILING DATE: 2001-02-12
PRIOR PILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 9218
SOFTWARE: SeqWin99, version 1.04
SEQ ID NO 8494
LENGTH: 209
 Query Match
Best Local Similarity
Matches 5; Conserv
 SEQ ID NO 15
 Sequence 15, Application US/11090916 Publication No. US20050260140A1
 GENERAL INFORMATION:
 Matches
 Query Match
Best Local Similarity
 APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
APPLICANT: Lee, Recolytic and Anti-Elastase Compounds and Methods of Use Thereof
 PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
 TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS FILE REFERENCE:
 SOFTWARE:
 TYPE: PRT
 TYPE: PRT
101 WCGPCR 106
 94 WCGPCR
 1 WCGPCK 6
 1 WCGPCK 6
 172
 5
 PatentIn version 3.3
 Conservative
 Conservative
 99
 Carl W.
 93.6%;
 93.6%; Score 44;
83.3%; Pred. No.
 Score 44; DB 7;
Pred. No. 1;
1; Mismatches
 Mismatches
 DB 6;
 0
 Length 209
 Length 172;
 Indels
 0
 0
 Gaps
 Gaps
 0
 0
```

RESULT 35 US-10-995-561-703 ; Sequence 703, Application US/10995561

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밁
 ; ORGANISM: Homo sapiens US-10-995-561-703
 CURRENT APPLICATION NUMBER: US/10/131,826A
CURRENT FILING DATE: 2002-04-24
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/05912
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/05912
PRIOR APPLICATION NUMBER: 60/059184
 US-10-131-826A-426
 RESULT 36
 Matches
 NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 703
 Sequence 426, Application US/10131826A Publication No. US20050245730A1
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 Publication No. US20050272054A1
 APPLICANT: CARGIIL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CARDITIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS
TITLE OF INVENTION: DETECTION AND USES THEREOF
FILE REFERENCE: CL001559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C128
 APPLICANT:
 APPLICANT: Baker, Kevin P.
 LENGTH: 126
TYPE: PRT
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICANT
 APPLICANT:
 APPLICANT:
 APPLICATION NUMBER: 60/059352
 APPLICATION NUMBER: 60/059263
FILING DATE: 1997-09-19
APPLICATION NUMBER: 60/059588
 62 WCGPCQ 67
 1 WCGPCK 6
 5; Conservative
 Smith, Victoria
 Godowski, Paul J.
Gurney, Austin L.
 Goddard, Audrey
 Gerritsen, Mary E.
 Wood,William
 Watanabe, Colin K
 Tumas,Daniel
 Stewart, Timothy A
 Sherwood, Steven
 Filvaroff, Ellen
 DeForge, Laura
 Zemin
 1997-09-18
 91.5%;
83.3%;
 Score 43;
Pred. No.
 Mismatches
 6;
 Length 126;
 Indels
 0
 Gaps
 쉱
 0
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 RESULT 38
US-10-821-234-860
 RESULT 37
US-10-467-962B-6
 US-10-131-826A-426
 US-10-467-962B-6
 SEQ ID NO 6
LENGTH: 183
TYPE: PRT
 SOFTWARE: pt
SEQ ID NO 860
LENGTH: 176
 Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 426
LENGTH: 747
TYPE: PRT
ORGANISM: Homo Sapien
 Sequence 860, Applic Publication No. US20 GENERAL INFORMATION:
 Query Match
Best Local &
 Matches
 Matches
 GENERAL INFORMATION:
 Sequence 6, Application US/1046 Publication No. US20050246784A1
 Query Match
Best Local Similarity
 FILE REFERENCE: 821A
CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILING DATE: 2004-04-07
CURRENT FILING DATE: US 60/462,047
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
 TITLE OF INVENTION: Identification of Herbicidally Active Substances FILE REFERENCE: 2000, 857
CURRENT APPLICATION NUMBER: US/10/467,962B
CURRENT FILING DATE: 2003-08-14
 PRIOR APPLICATION NUMBER: PCT/EP02/01466
PRIOR FILING DATE: 2002-02-13
NUMBER OF SEQ ID NOS: 109
 APPLICANT: Plesch,
APPLICANT: Blau, A
 TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
 APPLICANT:
 APPLICANT: Labat, Ivan
 SOFTWARE: PatentIn Vers. 2.0
 APPLICANT:
TYPE: PRT
ORGANISM: Homo sapiens
 ORGANISM: Arabidopsis thaliana
 Local Similarity hes 5; Conserv
 653 WCGPCQ 658
 105 WCGPC 109
 5;
 1 WCGPC 5
 1 WCGPCK 6
 Application US/10467962B
 Blau, Astrid
Daeschner, Klaus
Klein, Mathieu
 Andarmani, Su
Tang, Y. Tom
 Stache-Crain, Birgit
 Application US/10821234 o. US20050255114A1
 _SEQ_genes Version 1.0
 Conservative
 Conservative
 Gunnar
 100.0%;
 91.5%;
83.3%;
 Susan
 89.4%; Score 42;
100.0%; Pred. No.
 0
 1; Mismatches
 Score 43; DB 6; Pred. No. 3.8;
 Mismatches
 DB 6;
 Length 183;
 Length 747;
 Indels
 <u>.</u>
 <u>,</u>
 Gaps
 0,
 0,
```

Query Match

80.9%;

Score 38;

DB 6;

Length 176;

83.3%; Pred. No.

9

```
APPLICANT: Zhang, Jiem
APPLICANT: Zhou, Ping
APPLICANT: Zhou, Ping
APPLICANT: Cao, Yi-Cheng
APPLICANT: Cao, Yi-Cheng
APPLICANT: Cao, Yi-Cheng
APPLICANT: Darmanac, Radoje T.
TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides
FILE REFERENCE: 785CIP4CN
CURRENT APPLICATION NUMBER: US/11/000,463
CURRENT FILING DATE: 2004-11-29
PRIOR APPLICATION NUMBER: 10/291,265
PRIOR FILING DATE: 2002-11-08
PRIOR APPLICATION NUMBER: PT/US01/02623
PRIOR APPLICATION NUMBER: 09/92,279
PRIOR APPLICATION NUMBER: 09/92,279
PRIOR APPLICATION NUMBER: 09/92,279
PRIOR APPLICATION NUMBER: 09/91,404
PRIOR APPLICATION NUMBER: 09/491,404
PRIOR PILING DATE: 2000-01-25
PRIOR APPLICATION NUMBER: 09/617,746
PRIOR PILING DATE: 2000-07-17
PRIOR APPLICATION NUMBER: 09/631,451
PRIOR APPLICATION NUMBER: 09/631,451
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
 RESULT 39
US-11-000-463-344
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 Ś
 RESULT 40
US-10-878-556A-114
 S
 ; TYPE: PRT ; ORGANISM: Homo sapiens US-11-000-463-344
 В
FILE REFERENCE: 21762
CURRENT APPLICATION NUMBER: US/10/878,556A
CURRENT FILING DATE: 2004-06-28
NUMBER OF SEQ ID NOS: 199
SOFTWARE: PatentIn version 3.1
SEQ ID NO 114
LENGTH: 324
 Sequence 114, Application US/10878556A
Publication No. US20050266399A1
GENERAL INFORMATION:
APPLICANT: Hoffmann La-Roche Inc.
TITLE OF INVENTION: HCV regulated protein expression
 SOFTWARE: FastSEQ
SEQ ID NO 344
LENGTH: 269
 Query Match
Best Local Similarity
Matches 5; Conserv
 Best Local Similarity Matches 5; Conserv
 APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
 GENERAL INFORMATION
 Sequence 344, Publication No.
 NUMBER OF SEQ ID NOS: 944
COSTWARE: FastSEQ for Windows Version 3.0
 APPLICANT:
 APPLICANT: Tang, APPLICANT: Liu,
 PRIOR FILING DATE:
 APPLICANT:
 APPLICATION NUMBER: 09/633,870
 53 WCGHCK 58
 93 WCGHCK 98
 1 WCGPCK 6
 1 WCGPCK 6
 Qian, Xiaoho
Wang, Zhiwei
 Wehrman,
 Liu, Chenghua
Asundi, Vinod
 Application US/11000463
D. US20050266423A1
 Conservative
 Conservative
 Rui-hong
 Xiaohong
 Tom
 2000-09-15
 80.9%;
 ₽.
 0
 0,
 Score 38; DB
Pred. No. 12;
 Mismatches
 7;
 1;
 Length 269;
 Indels
 <u>.</u>
 ٥,
 Gaps
 Gaps
 0
 0
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Sequence 70, Application US/10523503

Publication No. US20060037102A1

GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
FILE REFERENCE: 16313-0236

CURRENT FILING DATE: 2005-02-02

PRIOR APPLICATION NUMBER: US/10/523,503

CURRENT FILING DATE: 2005-02-02

PRIOR APPLICATION NUMBER: US 60/400,803

PRIOR FILING DATE: 2002-08-02

NUMBER OF SEQ ID NOS: 163

SOFTWARE: Patentin version 3.2

SEQ ID NO 70

LENGTH: 361

TYPE: PRT

ORGANISM: Arabidopsis thaliana

US-10-523-503-70
 Sequence 1, Application US/10878556A
Publication No. US20050266399A1
GENERAL INFORMATION:
APPLICANT: Hoffmann La-Roche Inc.
TITLE OF INVENTION: HCV regulated protein expression
FILE REFERENCE: 21762
CURRENT APPLICATION NUMBER: US/10/878,556A
CURRENT FILING DATE: 2004-06-28
NUMBER OF SEQ ID NOS: 199
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
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 밁
 US-10-878-556A-114
 US-10-878-556A-1
 US-10-878-556A-1
 US-10-523-503-70
 RESULT 41
 Query Match
Best Local Similarity
Matches 5; Conserv
Query Match
Best Local Similarity
Matches 5; Conserv
 Matches
 Query Match
Best Local Similarity
 NAME/KEY: MISC FEATURE LOCATION: (307)..(307)
LOCATION: (307)..(307)
OTHER INFORMATION: Xaa is an undetermined amino acid residue PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: hsugp:006101-3-0
DATABASE ENTRY DATE: 2003-02-16
 ORGANISM: Homo sapiens
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: tr hum/q9bvh9
DATABASE ENTRY DATE: 2001-06-01
 TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
 TYPE: PRT
 LENGTH: 414
 108 WCGHCK 113
 51 WCGHCK 56
 1 WCGPCK 6
 1 WCGPCK 6
 5.
 Conservative
 Conservative
 Conservative
 80.9%;
 80.9%;
 80.9%;
Score 38; DB 6;
Pred. No. 16;
0; Mismatches
 0,
 Score 38;
Pred. No.
 <u>.</u>
 Score 38;
Pred. No.
 Mismatches
 Mismatches
 DB
13;
 DB
14;
 6.
 6,
 1;
 7:
 Length 414;
 Length 324;
 Length 361;
 Indels
 Indels
 <u>.</u>.
 0
 <u>,,</u>
 Gaps
 Gaps
 Gaps
 0
 0
 0,
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PRIOR APPLICATION UNMBER US 60/462,047
PRIOR FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
SOFTWARE: pt SEQ_genes Version 1.0
SEQ ID NO 1287
LENGTH: 505
TYPE: PRT
ORCANISM: Homo sapiens
US-10-821-234-1287
 PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILLING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
SOPTWARE: pt SEQ_genes Version 1.0
SEQ ID NO 1282
SEQ ID NO 1282
LENGTH: 440
TYPE: PRT
ORGANISM: Homo sapiens
US-10-821-234-1282
RESULT 45
US-10-821-234-1409
; Sequence 1409, Application US/10821234
 밁
 Query Match
Best Local Similarity
Marches 5; Conserve
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 US-10-821-234-1287
 US-10-821-234-1282
 Sequence 1282, Application US/10821234
Publication No. US20050255114A1
GENERAL INFORMATION:
APPLICANT: Labat, Ivan
APPLICANT: Stache-Crain, Birgit
 Sequence 1287, Application US/10821234 Publication No. US20050255114A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
Matches 5; Conserv
 FILE REFERENCE: 821A
CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILING DATE: 2004-04-07
 FILE REFERENCE: 821A
CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILING DATE: 2004-04-07
 APPLICANT: Andarmani, Susan
APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
 APPLICANT: Andarmani, Susan
APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
 APPLICANT: Labat, Ivan
APPLICANT: Stache-Crain,
 189 WCGHCK 194
 236
 1 WCGPCK 6
 1 WCGPCK 6
 WCCHCK 61
 WCGHCK 241
 Conservative
 Conservative
 80.9%;
 80.9%;
 Birgit
 0;
 Score 38;
Pred. No.
 Score 38; DB
Pred. No. 16;
 Mismatches
 DB 6;
18;
 <u>ه</u>.
 Length 505;
 Length 440
 0
 0,
 0,
 0
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RESULT 47
US-11-129-741-976
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 RESULT 46
US-10-895-064-976
 밁
 ; ORGANISM: Homo sapiens US-10-821-234-1409
 Sequence 976, Applicat
publication No. US2006
GENERAL INFORMATION:
APPLICANT: YUEN, KWOI
APPLICANT: WOO, CHI
 US-10-895-064-976
 APPLICANT: GUAN, YI
APPLICANT: CHAN, KWOK HUNG
APPLICANT: NICHOLLS, JOHN M.
APPLICANT: LEUNG, FREDERICK C.
TITLE OF INVENTION: THEREOR
TITLE OF INVENTION: THEREOR
FILE REFERENCE: V0690,0031
CURRENT APPLICATION NUMBER: US/10/895,064
CURRENT APPLICATION NUMBER: US/10/895,064
CURRENT FILING DATE: 2004-07-21
NUMBER OF SEQ ID NOS: 2918
SOFTWARE: Patentin version 3.2
SEQ ID NO 976
LENGTH: 32
TYPE: PRT
ORGANISM: COCONONAVICUS-HKU1
 Query Match
Best Local S
Matches 5
 APPLICANT: Stac...
APPLICANT: Andarmani, STAC...
APPLICANT: Tang, Y. Tor
 GENERAL INFORMATION:
APPLICANT: PEIRIS, JOSEPH S.M.
APPLICANT: YUEN, KWOK YUNG
APPLICANT: POON, LIT MAN
 SOFTWARE: pt_SEQ_genes Version 1.0 SEQ ID NO 1409
 Query Match
 Sequence 976, A Publication No.
 Matches
 Publication No. US20050255114A1
 Best Local Similarity 66.'
Matches 4; Conservative
 APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
FILE REFERENCE: 821A
CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
NUMBER OF SEQ ID NOS: 1704
 APPLICANT: Labat, Ivan
APPLICANT: Stache-Crain, Birgit
 TYPE: PRT
 ENGTH: 645
 Local Similarity
les 5; Conserv
 90 WCGHCK 95
 9
 1 WCGPCK 6
 1 WCGPCK 6
 Application US/11129741 o. US20060034853A1
 Application US/10895064
 Conservative
 ŪS20060018923A1
CHIU YAT PATRICK
KAR PUI SUSANNA
 76.6%;
66.7%;
 Susan
 Score 38;
Pred. No.
 Score 36; DB
Pred. No. 6.1;
 Mismatches
 DB 6;
 DB
 6
 Length 645;
 0;
 <u>.</u>
 Gaps
 Gaps
 0
 <u>,</u>
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APPLICANT: APPLICANT: APPLICANT:

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PRIOR APPLICATION NUMBER: 10/895,064
PRIOR FILING DATE: 2004-07-21
NUMBER OF SEQ ID NOS: 4257
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 3972
LENGTH: 32
TYPE: PRI
ORGANISM: COTONONAVITUS-HKU1
 밁
 Ś
 밁
 ; TYPE: PRT
; ORGANISM: Corononavirus-HKU1
US-11-129-741-976
 US-10-689-742-19
 US-11-129-741-3972
 RESULT 48
 US-11-129-741-3972
 CURRENT APPLICATION NUMBER: US/11/129,741
CURRENT FILING DATE: 2005-05-16
PRIOR APPLICATION NUMBER: 10/895,064
PRIOR FILING DATE: 2004-07-21
NUMBER OF SEQ ID NOS: 4257
SOFTWARE: PatentIn version 3.3
SEQ ID NO 976
 Sequence 19, Application US/10689742 Publication No. US20050250180A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
 Matches
 Query Match
 GENERAL INFORMATION:
 Publication No.
 Sequence 3972, Application US/11129741
 Matches
 APPLICANT: GUAN, YI
TITLE OF INVENTION: A NOVEL HUMAN VIRUS CAUSING RESPIRATORY TRACT
TITLE OF INVENTION: IMPECTION AND USES THEREOF
FILE REFERENCE: V0690.0044
CURRENT APPLICATION NUMBER: US/11/129,741
CURRENT FILING DATE: 2005-05-16
CURRENT FILING DATE: 2005-05-16
 APPLICANT: Jacobs, APPLICANT: McCoy, APPLICANT: LaValli
 APPLICANT: YUEN, KWOK YUNG
APPLICANT: WOO, CHIU YAT PATRICK
APPLICANT: LAU, KAR PUI SUSANNA
 TITLE OF INVENTION: A NOVEL HUMAN VIRUS CAUSING RESPIRATORY TRACT
TITLE OF INVENTION: INFECTION AND USES THEREOF
FILE REFERENCE: V0690.0044
 APPLICANT:
 APPLICANT:
 PPLICANT:
 LENGTH: 32
 Local Similarity
 1 WCGPCK 6
 9
 1 WCGPCK 6
 9 WCGSCR 14
 4;
 4;
McCoy, John M
LaVallie, Edward R
Racie, Lisa A
Evans, Cheryl
 WCGSCR 14
 PEIRIS, UN
 CHAN, KWOK HUI
POON, LIT MAN
 PEIRIS,
 Conservative
 Conservative
 US20060034853A1
 ĭ
 KWOK HUNG
 Kenneth
 JOSEPH S.M.
 JOSEPH S.M.
 76.6%;
66.7%;
 76.6%;
66.7%;
 Score 36; DB 7; Length 32; Pred. No. 6.1;
 Score 36;
Pred. No.
 Mismatches
 Mismatches
 DB 7;
 1; Indels
 Length 32;
 0;
 0,
 Gaps
 0
 0;
```

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APPLICANT: Treacy, Maurice
APPLICANT: Spaulding, Vikki
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POI
FILE REFERENCE: 00766.00091.10
CURRENT APPLICATION NUMBER: US/10/689,742
CURRENT FILING DATE: 2003-10-22
PRIOR APPLICATION NUMBER: 09/746,783
PRIOR APPLICATION NUMBER: 09/746,783
PRIOR FILING DATE: 2000-12-21
NUMBER OF SEQ ID NOS: 231
SOFTWARE: PatentIn version 3.2
SEQ ID NO 19
SEQ ID NO 19
 Query Match
Best Local Similarity
Tarches 5; Conserva
 RESULT 50
US-11-106-796-10
 밁
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 á
 ; ORGANISM: Homo sapiens
US-10-689-742-19
APPLICANT: Hastings, Craig
APPLICANT: Heeley, Robert
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
APPLICANT: Jung, Rudolf
APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOI
FILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
 RESULT 51
US-11-195-459-2
 US-11-106-796-10
 Sequence 10, Application US/11106796
Publication No. US20050282191A1
GENERAL INFORMATION:
 Sequence 2, Application US/11195459
Publication No. US20050278803A1
GENERAL INFORMATION:
 SEQ ID NO 10
LENGTH: 127
 Query Match
Best Local S
 FILE REFERENCE: UVMO:033US
CURRENT APPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2005-04-15
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
SOPTWARE: Patentin Ver. 2.1
 APPLICANT: Sewalt, Vincent
APPLICANT: Hastings, Crai
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
 APPLICANT: SUTOVSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
 ORGANISM: Homo sapiens
 TYPE: PRT
 LENGTH: 107
TYPE: PRT
 Local Similarity 100.0%;
les 5; Conservative 0;
 32 CGPCK 36
 32 CGPCK
 2 CGPCK 6
 2 CGPCK 6
 Conservative
 76.6%;
100.0%;
 <u>,</u>
 Score 36; DB 7; ; Pred. No. 15; 0; Mismatches
 Score 36; DB 6; pred. No. 13;
 Mismatches
 POLYNUCLEOTIDES ENCODING THEM
 FOR ALTERING THE DISULFIDE STATUS OF
 Length 127;
 Length 107;
 <u>.</u>.
 0
 0,
 0
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RESULT 53
US-11-072-512-2145
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 RESULT 52
US-11-195-459-4
 ; LENGTH: 128
; TYPE: PRT
; ORGANISM: Zea mays
US-11-195-459-2
 밁
 US-11-195-459-4
 Sequence 2145, Application US/11072512 Publication No. US20060029945A1 GENERAL INFORMATION:
 PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn version 3.0
SEQ ID NO 2
 Query Match
Best Local Similarity
Matches 4; Conser
 Matches
 Query Match
Best Local Similarity
 SEQ ID NO 4
LENGTH: 128
 Sequence 4, Application US/11195459 Publication No. US20050278803A1
 GENERAL INFORMATION:
 APPLICANT:
 APPLICANT: Jung, Rudolf
APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PROFILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-05
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR PELING DATE: 2001-12-03
PRIOR FILING DATE: 2001-12-03
PRIOR FILING DATE: 2001-12-03
 APPLICANT:
 APPLICANT:
 APPLICANT:
 PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
 APPLICANT: Sewalt, Vincent APPLICANT: Hastings, Crai
 SOFTWARE: PatentIn version 3.0
 APPLICANT:
 APPLICANT:
 APPLICANT
 APPLICANT
 APPLICANT
 TYPE: PRT
ORGANISM: Zea mays
 52 WCPPCR 57
 52 WCPPCR 57
 1 WCGPCK 6
 ISOGAI, TAKAO
SUGIYAMA, TOMOYASU
OTSUKI, TETSUJI
WAKAMATSU, AI
SATO, HIROYUKI
SATO, SHIZIKO
YAMAMOTO, JUN-ICHI
 1 WCGPCK 6
 4; Conservative
 4; Conservative
OTSUKA, KAORU
NAGAI, KEIICHI
IRIE, RYOTARO
 Meeley, Robert
Hantke, Sabine
 HIO,
 ISONO, YUUKO
HIO, YURI
 76.6%;
66.7%;
 76.6%;
66.7%;

 Mismatches

 Score 36;
Pred. No.
 Score 36;
Pred. No.
 Mismatches
 DB 7;
 Length 128,
 Length 128
 Indels
 Indels
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 Gaps
 Gaps
 0;
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 RESULT 54
US-10-510-386-126
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 Query Match
Best Local (
 Matches
 Sequence 126, Publication No.
 Matches
 APPLICANT:
APPLICANT:
APPLICANT:
 APPLICANT:
 TAMECHIKA,
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j,

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APPLICANT: NAGAHARI, KENJI
APPLICANT: NAGAHARI, KENJI
APPLICANT: MASUHO, YASUHIKO
ITILE OF INVENTION: NOVel full length cDNA
FILE REFERENCE: 084335-0191
CURRENT APPLICATION NUMBER: US/11/072,512
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: US 60/350,978
PRIOR APPLICATION NUMBER: US 60/350,978
PRIOR FILING DATE: 2002-01-25
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 4096
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 2145
LENGTH: 129
TYPER: PET
RESULT 55
US-11-054-281-58
; Sequence 58, Application US/11054281
; Publication No. US20060013813A1
; GENERAL INFORMATION:
 APPLICANT: Mezes et al.
; TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same
; FILE REFERENCE: 21402-240CIP
; CURRENT APPLICATION NUMBER: US/11/054,281
; CURRENT FILING DATE: 2005-02-08
; PRIOR APPLICATION NUMBER: 60/261,014
 ORGANISM: Bacillus licheniformis US-10-510-386-126
 ; TYPE: PRT; ORGANISM: Homo sapiens US-11-072-512-2145
 SOFTWARE: PatentIn version 3.3
SEQ ID NO 126
LENCTH: 163
TYPE: PRT
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 APPLICANT: Jorgensen, Steen Troels
APPLICANT: Olsen, Peter Bjarke
APPLICANT: Rasmussen, Michael Dolberg
TITLE OF INVENTION: Improved Bacillus Host Cell
FILE REFERENCE: 10294.204-US
CURRENT APPLICATION NUMBER: US/10/510,386
CURRENT FILING DATE: 2004-10-04
 APPLICANT: Andersen, Jens Tonne
APPLICANT: Clausen, Ib Groth
APPLICANT: Jorgensen, Steen Tr
 CURRENT FILING DATE: 2004-10-04 NUMBER OF SEQ ID NOS: 248
 Local Similarity 100.0%;
 62 WCPPCR 67
 74 CGPCK 78
 1 WCGPCK 6
 2 CGPCK 6
 5; Conservative
 SEKI, NAOHIKO
YOSHIKAWA, TSUTOMU
OTSUKA, MOTOYUKI
 Application US/10510386 o. US20050244922A1
 Conservative
 76.6%; Score 36; DB 6; 66.7%; Pred. No. 18;
 ICHIRO
 76.6%; Score 36; DB 7; 100.0%; Pred. No. 15;
 0; Mismatches
 Mismatches
 1; Indels
 0
 Length 163;
 Length 129
 Indels
 0
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 Gaps
 Gaps
 0
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밁
 Query Match
Best Local Similarity
Watches 5; Conserv
 S
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 S
 US-11-054-281-59
 US-11-054-281-59
 RESULT 56
 ; ORGANISM: Mus musculus
US-11-054-281-58
 NUMBER OF SEQ ID NOS: 3
SOFTWARE: PatentIn Ver.
SEQ ID NO 59
 GENERAL INFORMATION:
 Sequence 59, Application US/11054281
Publication No. US20060013813A1
 Matches
 Query Match
Best Local 9
 SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 58
LENGTH: 354
 PRIOR APPLICATION NUMBER: 60/261,018
PRIOR TILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/318,410
PRIOR FILING DATE: 2001-09-10
PRIOR APPLICATION NUMBER: 60/261,013
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/261,026
 CURRENT APPLICATION NUMBER: US/11/054,281
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: 60/261,014
PRIOR FILING DATE: 2001-01-11
 APPLICANT: Mezes et al. TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same FILE REFERENCE: 21402-240CIP
 PRIOR APPLICATION NUMBER: 60/261,013
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/261,026
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/261,029
PRIOR FILING DATE: 2001-01-11
 PRIOR FILING DATE: 2001-08-17
PRIOR APPLICATION NUMBER: 10/044,564
PRIOR FILING DATE: 2002-01-11
 PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/261,029
 PRIOR APPLICATION NUMBER: 10/044,564 PRIOR FILING DATE: 2002-01-11
 PRIOR APPLICATION NUMBER: 60/313,170 PRIOR FILING DATE: 2001-08-17
 NUMBER OF SEQ ID NOS: 324
 TYPE: PRT ORGANISM: Rattus norvegicus
 TYPE: PRT
 LENGTH:
 FILING DATE:
 APPLICATION NUMBER: 60/318,410 FILING DATE: 2001-09-10
 APPLICATION NUMBER: 60/313,170
 FILING DATE:
 APPLICATION NUMBER: 60/261,018
 FILING DATE: 2001-01-11
 144 CGPCK 148
163 CGPCK 167
 2 CGPCK 6
 2 CGPCK 6
 Similarity 5; Conserv
 Conservative
 Conservative
 2001-01-1
 2002-01-11
 2001-01-11
 76.6%;
100.0%;
 76.6%; Score 36;
100.0%; Pred. No.
cive 0; Mismatc
 Score 36; DB 7;
; Pred. No. 29;
 0;
 Mismatches
 Mismatches
 DB 7;
30;
 ٥,
 Length 354;
 Length 373;
 Indels
 Indels
 0
 0
 Gaps
 0
 0
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US-11-090-439-54
Sequence 54, Application US/11090439
Publication No. US20050266442A1
GENERAL INFORMATION:
APPLICANT: Squillace, Rachel
APPLICANT: Weiner, Michael P.
APPLICANT: Weiner, Michael P.
TITLE OF INVENTION: Amgiomyolipoma Cell and Method of Use Thereof
FILE REFERENCE: 24318-502
FILE REFERENCE: 24318-502
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 RESULT 57
US-11-090-439-52
 US-11-054-281-60
 US-11-090-439-54
 RESULT 58
 US-11-090-439-52
 PRIOR APPLICATION NUMBER: 60/556,344
PRIOR FILING DATE: 2004-03-25
NUMBER OF SEQ ID NOS: 62
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 52
 Sequence 52, Application US/11090439
Publication No. US20050286442A1
GENERAL INFORMATION:
APPLICANT: Squillace, Rachel
APPLICANT: Weiner, Michael P.
TITLE OF INVENTION: Immorttalized Human Tuberous Sclerosis Null
TITLE OF INVENTION: Augiomyolipoma Cell and Method of Use Thereof
FILE REFERENCE: 24318-502
 Sequence 60, Application US/11054281 Publication No. US20060013813A1 GENERAL INFORMATION:
 CURRENT APPLICATION NUMBER: US/11/090,439
CURRENT FILING DATE: 2005-03-25
PRIOR APPLICATION NUMBER: 60/556,344
PRIOR FILING DATE: 2004-03-25
NUMBER OF SEQ ID NOS: 62
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 54
 Query Match
Best Local Similarity
 Best Local Similarity Matches 5; Conserv
 Query Match
 Matches
 APPLICANT: Mezes et al.
TITLE OF INVENTION: Proteins and Nucleic Acids Encoding
FILE REFERENCE: 21402-240CIP
 CURRENT APPLICATION NUMBER: US/11/090,439
CURRENT FILING DATE: 2005-03-25
CURRENT APPLICATION NUMBER: US/11/054,281
 LENGTH: 380
TYPE: PRT
ORGANISM: Homo sapiens
 ORGANISM: Homo sapiens
 TYPE: PRT
 LENGTH: 380
 170 CGPCK 174
 170 CGPCK 174
 2 CGPCK 6
 2 CGPCK 6
 <u>ن</u>
 Conservative
 Conservative
 76.6%; but
100.0%; Pre
 76.6%; Score 36; DB 7; 100.0%; Pred. No. 31;
 0,
 Score 36;
Pred. No.
 Mismatches
 Mismatches
 DB 7;
31;
 <u>.</u>
 0
 Length 380,
 Length 380;
 Indels
 <u>.</u>.
 <u>.</u>
 Gaps
 Gaps
 0
 0
```

CURRENT FILING DATE:

2005-02-08

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RESULT 60
US-10-763-712A-10
US-10-763-712A

PUBLICANT INFORMATION:
APPLICANT: Solazyme, Inc.
APPLICANT: Dillon, Harrison F.
TITLE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen
TITLE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen
TITLE OF INVENTION: Workering Microbial Hydrogen
CURRENT APPLICATION NUMBER: US/10/763,712A
CURRENT FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: US 10/287,750
PRIOR APPLICATION NUMBER: US 10/287,750
PRIOR APPLICATION NUMBER: US 10/411,910
PRIOR APPLICATION NUMBER: US 003-04-12
PRIOR PILING DATE: 2003-04-12
PRIOR PILING DATE: 2003-09-03

**THATTER OF SEA TO NOS: 184
 SOPTWARE: Patentin Ver. 2
SEQ ID NO 60
LENGTH: 380
TYPE: PRT
ORGANISM: Homo sapiens
US-11-054-281-60
RESULT 61
US-10-763-712A-11
Sequence 11, Application US/10763712A
; Publication No. US20050266541A1
 S
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 Query Match
Best Local Similarity
Marches 5; Conserve
 밁
 US-10-763-712A-10
 PRIOR APPLICATION NUMBER: 60/318,410
PRIOR FILING DATE: 2001-09-10
PRIOR APPLICATION NUMBER: 60/261,013
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/261,026
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/261,029
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/313,170
PRIOR APPLICATION NUMBER: 60/313,170
PRIOR FILING DATE: 2001-08-17
PRIOR APPLICATION NUMBER: 10/044,564
PRIOR FILING DATE: 2002-01-11
NUMBER OF SEQ ID NOS: 324
PRIOR FILING DATE: 2002-01-11
NUMBER OF SEQ ID NOS: 324
 Query Match
Best Local Similarity
Matches 5; Conserv
 NUMBER OF SEQ ID NOS: 184
SOFTWARE: PatentIn version
SEQ ID NO 10
 PRIOR APPLICATION NUMBER: 60/261,018 PRIOR FILING DATE: 2001-01-11
 PRIOR APPLICATION NUMBER: 60/261,014 PRIOR FILING DATE: 2001-01-11
 TYPE: PRT ORGANISM: Clostridium perfringens
 LENGTH: 572
 170 CGPCK 174
 86
 2 CGPCK 6
 2 CGPCK 6
 CGPCK 102
 Conservative
 Conservative
 version 3.2
 76.6%; Score 36; DB 7; 100.0%; Pred. No. 31;
 76.6%; Score 36; DB 6; Length 572; 100.0%; Pred. No. 40;
 0; Mismatches
 0; Mismatches
 0
 0
 Length 380;
 Indels
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 Gaps
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Solazyme, Inc. Dillon, Harrison F.

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Sequence 37, Application US/10763712A

Publication No. US20050266541A1

GENERAL INFORMATION:
APPLICANT: Solazyme, Inc.

APPLICANT: Dillon, Harrison F.

ITILE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen

ITILE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen

ITILE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen

ITILE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen

ITILE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen

ITILE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen

ITILE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen

ITILE OF INVENTION: MOMBER: US/10/763,712A

CURRENT APPLICATION NUMBER: US 10/287,750

PRIOR APPLICATION NUMBER: US 10/411,910

PRIOR APPLICATION NUMBER: US 10/411,910

PRIOR APPLICATION NUMBER: US 60/500,032

PRIOR APPLICATION NUMBER: US 60/500,032

PRIOR FILING DATE: 2003-09-03

NUMBER OF SEQ ID NOS: 184
 RESULT 63
US-10-763-712A-109
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 S
 RESULT 62
US-10-763-712A-37
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 US-10-763-712A-37
 US-10-763-712A-11
Sequence 109, Application US/10763712A Publication No. US20050266541A1 GENERAL INFORMATION:
APPLICANT: Solazyme, Inc.
APPLICANT: Dillon, Harrison F.
 SOFTWARE: PatentIn version 3.2 SEQ ID NO 37
 Query Match 76.6%;
Best Local Similarity 100.0%;
 SOFTWARE: PatentIn version 3.2 SEQ ID NO 11
 GENERAL INFORMATION:
 Matches
 Matches
 Query Match 76.6%; Score 36; DB 6; Best Local Similarity 100.0%; Pred. No. 40;
 APPLICANT: Solazyme, Inc.
APPLICANT: Solazyme, Inc.
APPLICANT: Dillon, Harrison F.
APPLICANT: Dillon, Harrison F.
TITLE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen
TITLE OF INVENTION: Production
FILE REFERENCE: H2042101-CIP
CURRENT APPLICATION NUMBER: US/10/763,712A
CURRENT FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: US 10/287,750
PRIOR PILING DATE: 2002-11-04
PRIOR PILING DATE: 2003-04-12
PRIOR APPLICATION NUMBER: US 10/411,910
PRIOR APPLICATION NUMBER: US 10/411,910
PRIOR PILING DATE: 2003-09-03
PRIOR FILING DATE: 2003-09-03
PRIOR FILING DATE: 2003-09-03
PRIOR FILING DATE: 2003-09-03
PRIOR FILING DATE: 2003-09-03
 LENGTH: 572
TYPE: PRT
ORGANISM: Clostridium perfringens
 TYPE: PRT ORGANISM: Clostridium perfringens
 LENGTH: 572
 98 CGPCK 102
 98 CGPCK 102
 2 CGPCK 6
 2 CGPCK 6
 5; Conservative
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 Conservative
 76.6%; Score 36; DB 6; 100.0%; Pred. No. 40;
 0; Mismatches
 0; Mismatches
 0,
 ٥,
 Length 572;
 Length 572;
 Indels
 0
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 Gaps
 Gaps
 0
 0
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 SOFTWARE: PatentIn version 3.2
SEQ ID NO 109
LENGTH: 572
TYPE: PRT
ORGANISM: C. perfringens
US-10-763-712A-109
 ; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 110
; LENGTH: 572
; TYPE: PRT
; ORGANISM: C. perfringens
US-10-763-712A-110
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 US-10-055-877-255
 RESULT 65
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 US-10-763-712A-110
 RESULT 64
 CURRENT APPLICATION NUMBER: US/10/763,712A
CURRENT FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: US 10/287,750
PRIOR FILING DATE: 2002-11-04
PRIOR APPLICATION NUMBER: US 10/411,910
PRIOR APPLICATION NUMBER: US 10/411,910
PRIOR FILING DATE: 2003-04-12
PRIOR APPLICATION NUMBER: US 60/500,032
PRIOR FILING DATE: 2003-09-03
NUMBER OF SEQ ID NOS: 184
 Sequence 255, Application US/1005877
Publication No. US20050288241A1
GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
APPLICANT: Tcherney, Velizar
 Query Match
Best Local 9
 GENERAL INFORMATION:
 Sequence 110, Application US/10763712A Publication No. US20050266541A1
 Matches
 Best
 Query Match
 Matches
 FILE REFERENCE: H2042101-CIP
CURRENT APPLICATION NUMBER: US/10/763,712A
CURRENT FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: US 10/287,750
PRIOR FILING DATE: 2002-11-04
PRIOR APPLICATION NUMBER: US 10/411,910
PRIOR FILING DATE: 2003-04-12
 PRIOR APPLICATION NUMBER: US 60/500,032
PRIOR FILING DATE: 2003-09-03
NUMBER OF SEQ ID NOS: 184
 APPLICANT: Solazyme, Inc.
APPLICANT: Dillon, Harrison F.
TITLE OF INVENTION: Methods and
TITLE OF INVENTION: Production
 TITLE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen TITLE OF INVENTION: Production
 APPLICANT:
 FILE REFERENCE: H2042101-CIP
 Local
 98 CGPCK 102
 98 CGPCK 102
 l Similarity
5; Conserva
 2 CGPCK 6
 2 CGPCK 6
 h 76.6%; Somilarity 100.0%; I 5; Conservative 0;
 Conservative
 100.0%;
 76.6%;
 score 36; DB; Pred. No. 40; 0; Mismatches
 Score 36; pred. No.
 0,
 Compositions for Evolving Microbial Hydrogen
 Mismatches
 DB 6;
 DB 6;
 0
 0;
 Length 572;
 Length 572;
 Indels
 Indels
 0,
 ٥,
 0
 0,
 ; SEQ ID NO 255
, LENGTH: 654
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-055-877-255
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 US-10-055-877-254
 RESULT 66
 CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION UNMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR FILING DATE: 2001-01-23,598
PRIOR FILING DATE: 2001-01-23
PRIOR PELICATION NUMBER: 60/263,799
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR FILING DATE: 2001-01-24
PRIOR FILING DATE: 2001-01-24
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR APPLICATION NUMBER: 60/264,139
Sequence 254, Application US/100558
Publication No. US20050288241A1
GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
 Query Match
Best Local Similarity
Matches 5; Conserv
 APPLICANT:
APPLICANT:
APPLICANT:
 APPLICANT:
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APPLICANT:
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APPLICANT:
APPLICANT:
 Remaining Prior Application
NUMBER OF SEQ ID NOS: 512
SOFTWARE: PatentIn Ver. 2.1
 PRIOR APPLICATION NUMBER: 60/275,990 PRIOR FILING DATE: 2001-03-14
 APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
 APPLICANT:
 APPLICANT:
APPLICANT:
 PRIOR APPLICATION NUMBER: 60/275,927 PRIOR FILING DATE: 2001-03-14
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: 60/272,870
 PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,478
 PRIOR APPLICATION NUMBER: 60/263,351
 PRIOR FILING DATE: 2001-01-26
 APPLICANT:
 APPLICANT:
 PPLICANT:
 PPLICANT:
 PPLICANT
 PPLICANT
 PPLICANT:
 FILING DATE: 2001-03-02
 247 CGPCK 251
 2 CGPCK 6
 Kekuda, Ramesh
Guo, Xiaojia
Zerhusen, Bryan
 Vernet, Corine
Taupier Jr., Raymond
Pena, Carol
 Gerlach, Valerie
Spytek, Kimberly
 Shenoy, Suresh
 Gusev, Vladimir
 Baumgartner, Jason
Shimkets, Richard
 Patturajan, Meera
Burgess, Cahterine
 Andrew, Davio
Mezes, Peter
 Ballinger,
 Anderson, David
 Eisen, Andrew
 Ratelli, Luca
 Application US/10055877
o. US20050288241A1
 Wolenc, Adam
 Conservative
 David
 valerie
 76.6%; Sur
100.0%; Pr
 data removed - See File Wrapper or PALM
 Score 36; pred. No.
 Mismatches
 DB 6;
43;
 0
 Length 654;
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Gaps

0

Miller, Charles

Velizar

Tchernev, Zhong, Mei

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RESULT 67
US-10-831-997-4
 ; LENGTH: 660
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-055-877-254
Sequence 4, Application US/10831997
Publication No. US20050244834A1
 Query Match
Best Local &
 Remaining Prior Application NUMBER OF SEQ ID NOS: 512 SOFTWARE: PatentIn Ver. 2 SEQ ID NO 254
 Matches
 CURRENT APPLICATION NUMBER: US/10/055,877
 PRIOR APPLICATION NUMBER: 60/262,892 PRIOR FILING DATE: 2001-01-19
 RIOR FILING DATE:
 RIOR APPLICATION NUMBER: 60/263,598
 PHICANT: Boldog, Ference
TILE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
 Local Similarity nes 5; Conserv
 APPLICATION NUMBER: 60/263,799 FILING DATE: 2001-01-24
 APPLICATION NUMBER: 60/264,117 FILING DATE: 2001-01-25
 FILING DATE: 2001-03-1
 APPLICATION NUMBER: 60/275,990
 FILING DATE
 APPLICATION NUMBER: 60/264,139
 APPLICATION NUMBER: 60/275,927
 APPLICATION NUMBER: 60/272,870
 APPLICATION NUMBER: 60/263,351
 APPLICATION NUMBER: 60/264,478
 249 CGPCK 253
 LING DATE:
 2 CGPCK 6
 Anderson, David
Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
 Vernet, Corine
Taupier Jr., Raymond
Pena, Carol
 Shenoy, Suresh
 Mezes, Peter
Patturajan, Meera
Burgess, Cahterine
 Ratelli, Luca
Kekuda, Ramesh
 Casman, Stacie
 Gusev, Vladimir
 Baumgartner, Jason
Shimkets, Richard
 Burgess, Cahte
Eisen, Andrew
 Zerhusen, Bryan
Andrew, David
 Conservative
 Application data removed - See File Wrapper or PALM
 Xiaojia
 2001-01-25
 2001-01-23
 2001-03-02
 2001-01-30
 2001-01-26
 2002-01-22
 76.6%;
 0
 Score 36; DB 6; Pred. No. 44;
 Mismatches
 0; Indels
 Length 660;
 Gaps
 0
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 SOPTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 961
TYPE: PRT
ORGANISM: Homo sapiens
US-10-831-997-4
 US-10-055-877-252
 Sequence 252, Application US/10055877 Publication No. US20050288241A1
 Matches
 Query Match
Best Local
 GENERAL INFORMATION:
 APPLICANT: Daley, George Q.
APPLICANT: McCarthy, Jeanette J.
APPLICANT: S825.1027-001
CURRENT FILING DATE: 2825.1027-001
CURRENT FILING DATE: 2004-04-26
PRIOR APPLICATION NUMBER: US/09/657,472
PRIOR FILING DATE: 2000-09-07
PRIOR APPLICATION NUMBER: US 60/153,357
PRIOR APPLICATION NUMBER: US 60/220,947
PRIOR APPLICATION NUMBER: US 60/220,947
 PRIOR FILING DATE: 2000-07-26
PRIOR APPLICATION NUMBER: US 60/225,724
PRIOR FILING DATE: 2000-08-16
NUMBER OF SEQ ID NOS: 2551
 APPLICANT: Lander, Eric S.
APPLICANT: Cargill, Michele
APPLICANT: Ireland, James S.
APPLICANT: Bolk, Stacey
APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
 APPLICANT:
 APPLICANT:
 Local Similarity 100.0%; nes 5; Conservative (
 403 CGPCK 407
 2 CGPCK 6
 Guo, Xiaojia
Zerhusen, Bryan
Andrew, David
Mezes, Peter
Patturajan, Meera
Burgess, Cahterine
 DeCristofaro, Marc
 Vernet, Corine
Taupier Jr., Raymond
Pena, Carol
 Baumgartner, Jason
Shimkets, Richard
Gusev, Vladimir
 Anderson, David
Ballinger, Rober
 Padigaru, Muralidhara
Miller, Charles
 Casman, Stacie
 Shenoy, Suresh
 Eisen, Andrew
Wolenc, Adam
 Ratelli, Luca
Kekuda, Ramesh
 Spytek, Kimberly
 Gerlach,
 Zhong, Mei
 Ichernev,
 Valerie
 Velizar
 DB 6;
 Length 961;
 Indels
 0;
 Gaps
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0

CURRENT APPLICATION NUMBER: US/10/055,877

CURRENT FILING DATE:

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 US-10-055-877-73
 RESULT 69
 ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-055-877-252
 SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 252
LENGTH: 1857
 GENERAL INFORMATION:
 Sequence 73, Application US/10055877 Publication No. US20050288241A1
 Query Match
Best Local (
 Matches
 APPLICANT:
 PRIOR
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 APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
 Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 512
 APPLICANT:
 APPLICANT:
 APPLICANT:
 PRIOR FILING DATE:
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 APPLICATION NUMBER: 60/263,799
FILING DATE: 2001-01-23
 APPLICATION NUMBER: 60/272,870
 NT FILING DATE: 2002-01-22
APPLICATION NUMBER: 60/262,892
 FILING DATE: 2001-01-25
 APPLICATION NUMBER: 60/264,117
 APPLICATION NUMBER: 60/263,598
 APPLICATION NUMBER: 60/264,478
 APPLICATION NUMBER: 60/264,139
 APPLICATION NUMBER: 60/275,927
 FILING DATE:
 APPLICATION NUMBER: 60/275,990
 FILING DATE:
 FILING DATE:
 APPLICATION NUMBER: 60/263,351
 FILING DATE:
 FILING DATE:
 65 CGPCK 69
 2 CGPCK 6
 Similarity 5; Conserv
 Guo, Xiaojia
Zerhusen, Bryan
Andrew, David
 Vernet, Corine
Taupier Jr., Raymond
Pena, Carol
 Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
 Padigaru, Murazi
Miller, Charles
Boldog, Ference
 Casman, Stacie
 Shenoy, Suresh
 Gusev, Vladimir
 Wolenc, Adam
Baumgartner, Jason
Shimkets, Richard
 Eisen, Andrew
 Patturajan, Meera
Burgess, Cahterine
 Mezes, Peter
 Ratelli, Luca
Kekuda, Ramesh
 Anderson, David
 Tchernev,
 Conservative
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 2001-03-14
 2001-03-14
 2001-03-02
 2001-01-30
 2001-01-26
 2001-01-25
 2001-01-19
 76.6%;
100.0%;
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 Robert
 0;
 Score 36;
Pred. No.
 Mismatches
 .
B6;
 Length 1857;
 Indels
 0
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밁
 ; TYPE: PRT; ORGANISM: Homo sapiens US-10-055-877-73
 US-10-055-877-253
 RESULT 70
 PRIOR FILING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR FILING DATE: 2001-01-24
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
 Query Match
Best Local S
Matches 5
 SOFTWARE: POSEQ ID NO 73
 GENERAL INFORMATION:
 Sequence 253, P
Publication No.
 APPLICANT:
 PRIOR FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: 60/272,870
 FILE REFERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
 APPLICANT:
APPLICANT:
 APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
 Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 512
 PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/275,927
 TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby FILE REFERENCE: 21402-251
 PRIOR FILING DATE: 2001-03-14
 PRIOR FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: 60/275,990
 APPLICANT:
 APPLICANT:
 APPLICANT:
 PRIOR APPLICATION NUMBER: 60/264,478
 APPLICANT
 APPLICANT
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 APPLICANT
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 APPLICANT:
 APPLICANT:
APPLICANT:
 APPLICANT
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 APPLICANT
 LENGTH: 2084
 317 CGPCK 321
 2 CGPCK 6
 5
 Similarity
 PatentIn Ver. 2.1
 Wolenc, Adam
Baumgattner, Jason
Shimkets, Richard
Gusev, Vladimir
Vernet, Corine
Taupier Jr., Raymond
 Zerhusen, L.,
Zerhusen, L.,
Zew, David
 Mezes, Peter
Patturajan, Meera
Burgess, Cahterine
 Guo, Xiaojia
Zerhusen, Bryan
 Gerlach, Valerie
Spytek, Kimberly
 Eisen, Andrew
 Ratelli, Luca
Kekuda, Ramesh
 Ballinger,
 Anderson, David
 Zhong,
 Tchernev,
 Pena, Carol
 Application US/10055877
o. US20050288241A1
 Conservative
 Mer
 Velizar
 76.6%; Score 36; 100.0%; Pred. No.
 0;
 Mismatches
 DB 6;
92;
 Length 2084;
 Indels
 ٥,
 Gaps
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APPLICANT: Li, Li
APPLICANT: Casman, Stacie
FILE REFERENCE: 21402-21
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT APPLICATION NUMBER: 00/262,892
FRIOR FILING DATE: 2001-01-29
FRIOR PLILNG DATE: 2001-01-23
FRIOR FILING DATE: 2001-01-24
FRIOR FILING DATE: 2001-01-24
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FRIOR PLILNG DATE: 2001-01-25
FRIOR APPLICATION NUMBER: 60/264,478
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FRIOR APPLICATION NUMBER: 60/264,478
FRIOR PLILNG DATE: 2001-01-36
FRIOR PLILNG DATE: 2001-03-02
FRIOR PLILNG DATE: 2001-03-04
FRIOR PLING DATE: 2001-03-04
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FRIOR PLING DATE: 2001-03-04
FRIOR PLING
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Query Match 76.6%; Score 36; DB 6; Length 2098; Best Local Similarity 100.0%; Pred. No. 93; Matches 5; Conservative 0; Mismatches 0; Indels 0;

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Gaps

2 CGPCK 6

Db 267 CGPCK 271

Search completed: February 23, 2006, 00:48:12 Job time : 15.5 secs

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